

International Nonproprietary Names for Pharmaceutical Substances (INN)

RECOMMENDED International Nonproprietary Names: List 68

Notice is hereby given that, in accordance with paragraph 7 of the Procedure for the Selection of Recommended International Nonproprietary Names for Pharmaceutical Substances [*Off. Rec. Wld Health Org.*, 1955, **60**, 3 (Resolution EB15.R7); 1969, **173**, 10 (Resolution EB43.R9); Resolution EB115.R4 (EB115/2005/REC/1)], the following names are selected as Recommended International Nonproprietary Names. The inclusion of a name in the lists of Recommended International Nonproprietary Names does not imply any recommendation of the use of the substance in medicine or pharmacy.

Lists of Proposed (1–105) and Recommended (1–66) International Nonproprietary Names can be found in *Cumulative List No. 14, 2011* (available in CD-ROM only).

Dénominations communes internationales des Substances pharmaceutiques (DCI)

Dénominations communes internationales RECOMMANDÉES: Liste 68

Il est notifié que, conformément aux dispositions du paragraphe 7 de la Procédure à suivre en vue du choix de Dénominations communes internationales recommandées pour les Substances pharmaceutiques [*Actes off. Org. mond. Santé*, 1955, **60**, 3 (résolution EB15.R7); 1969, **173**, 10 (résolution EB43.R9); résolution EB115.R4 (EB115/2005/REC/1)] les dénominations ci-dessous sont choisies par l'Organisation mondiale de la Santé en tant que dénominations communes internationales recommandées. L'inclusion d'une dénomination dans les listes de DCI recommandées n'implique aucune recommandation en vue de l'utilisation de la substance correspondante en médecine ou en pharmacie.

On trouvera d'autres listes de Dénominations communes internationales proposées (1–105) et recommandées (1–66) dans la *Liste récapitulative No. 14, 2011* (disponible sur CD-ROM seulement).

Denominaciones Comunes Internacionales para las Sustancias Farmacéuticas (DCI)

Denominaciones Comunes Internacionales RECOMENDADAS: Lista 68

De conformidad con lo que dispone el párrafo 7 del Procedimiento de Selección de Denominaciones Comunes Internacionales Recomendadas para las Sustancias Farmacéuticas [Act. Of. Mund. Salud, 1955, **60**, 3 (Resolución EB15.R7); 1969, **173**, 10 (Resolución EB43.R9); Resolución EB115.R4 (EB115/2005/REC/1) EB115.R4 (EB115/2005/REC/1)], se comunica por el presente anuncio que las denominaciones que a continuación se expresan han sido seleccionadas como Denominaciones Comunes Internacionales Recomendadas. La inclusión de una denominación en las listas de las Denominaciones Comunes Recomendadas no supone recomendación alguna en favor del empleo de la sustancia respectiva en medicina o en farmacia.

Las listas de Denominaciones Comunes Internacionales Propuestas (1–105) y Recomendadas (1–66) se encuentran reunidas en *Cumulative List No. 14, 2011* (disponible sólo en CD-ROM).

Latin, English, French, Spanish:
Recommended INN

Chemical name or description; Molecular formula; Graphic formula

DCI Recommandée

Nom chimique ou description; Formule brute; Formule développée

DCI Recomendada

Nombre químico o descripción; Fórmula molecular; Fórmula desarrollada

acidum deoxycholicum
deoxycholic acid

3 α ,12 α -dihydroxy-5 β -cholan-24-oic acid

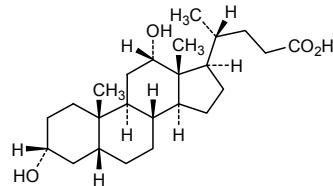
acide désoxycholique

acide 3 α ,12 α -dihydroxy-5 β -cholan-24-oïque

ácido desoxicólico

ácido 3 α ,12 α -dihidroxi-5 β -colan-24-oíco

C₂₄H₄₀O₄



acidum florilglutamicum (¹⁸F)
florilglutamic acid (¹⁸F)

(4S)-4-(3-[¹⁸F]fluoropropyl)-L-glutamic acid

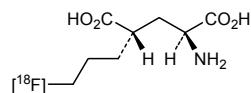
acide florilglutamique (¹⁸F)

acide (4S)-4-(3-[¹⁸F]fluoropropyl)-L-glutamique

ácido florilglútamico (¹⁸F)

ácido (4S)-4-(3-[¹⁸F]fluoropropil)-L-glutámico

C₈H₁₄¹⁸FNO₄



acidum tiazoticum
tiazotic acid

[(5-methyl-1*H*-1,2,4-triazol-3-yl)sulfanyl]acetic acid

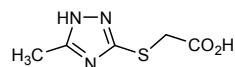
acide tiazotique

acide [(5-méthyl-1*H*-1,2,4-triazol-3-yl)sulfanyl]acétique

ácido tiazótico

ácido [(5-metil-1*H*-1,2,4-triazol-3-il)sulfanil]acético

C₅H₇N₃O₂S



brexpiprazolum
brexpiprazole

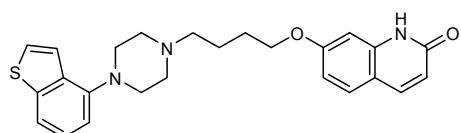
7-[4-[4-(1-benzothiophen-4-yl)piperazin-1-yl]butoxy]quinolin-2(1H)-one

brexpiprazole

7-[4-[4-(1-benzothiophén-4-yl)pipérazin-1-yl]butoxy]quinoléin-2(1H)-one

brexpiprazol

7-[4-[4-(1-benzotiofen-4-il)piperazin-1-il]butoxi]quinolin-2(1H)-ona

C₂₅H₂₇N₃O₂S**buparlisibum**
buparlisib

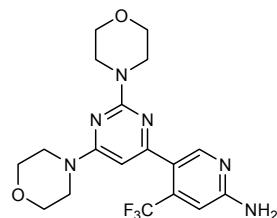
5-[2,6-bis(morpholin-4-yl)pyrimidin-4-yl]-4-(trifluoromethyl)pyridin-2-amine

buparlisib

5-[2,6-bis(morpholin-4-yl)pyrimidin-4-yl]-4-(trifluorométhyl)pyridin-2-amine

buparlisib

5-[2,6-bis(morfolin-4-il)pirimidin-4-il]-4-(trifluorometil)piridin-2-amina

C₁₈H₂₁F₃N₆O₂**camicinalum**
camicinal

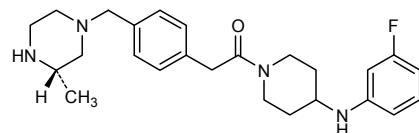
1-{4-[(3-fluorophenyl)amino]piperidin-1-yl}-2-(4-[(3S)-3-methylpiperazin-1-yl]methyl)ethan-1-one

camicinal

1-{4-[(3-fluorophényl)amino]pipéridin-1-yl}-2-(4-[(3S)-3-méthylpipérazin-1-yl]méthyl)phényléthan-1-one

camicinal

1-{4-[(3-fluorofenil)amino]piperidin-1-il}-2-(4-[(3S)-3-metilpiperazin-1-il]metil)fenil)etan-1-ona

C₂₅H₃₃FN₄O

caplacizumabum #
caplacizumab

immunoglobulin VH-linker-VH fragment, anti-[*Homo sapiens* VWF (von Willebrand factor) A1 domain], humanized monoclonal antibody;
 VH-linker-VH chain (1-259) [humanized VH (*Homo sapiens* IGHV3-23*04 (82.50%) -(IGHD)-IGHJ4*01 L123>Q (123) [8.8.21] (1-128)] -trialanyl linker (129-131) -[humanized VH (*Homo sapiens* IGHV3-23*04 (82.50%) -(IGHD)-IGHJ4*01 L123>Q (254) [8.8.21] (132-259)

caplacizumab

immunoglobuline fragment VH-linker-VH, anti-[*Homo sapiens* VWF (facteur de von Willebrand) domaine A1], anticorps monoclonal humanisé;
 chaîne VH-linker-VH (1-259) [VH humanisé (*Homo sapiens* IGHV3-23*04 (82.50%) -(IGHD)-IGHJ4*01 L123>Q (123) [8.8.21] (1-128)] -trialanyl linker (129-131) -[VH humanisé (*Homo sapiens* IGHV3-23*04 (82.50%) -(IGHD)-IGHJ4*01 L123>Q (254) [8.8.21] (132-259)

caplacizumab

inmunoglobulina fragmento VH-conector-VH, anti-[VWF (factor de von Willebrand) de *Homo sapiens* dominio A1], anticuerpo monoclonal humanizado;
 cadena VH-conector-VH (1-259) [VH humanizado (*Homo sapiens* IGHV3-23*04 (82.50%) -(IGHD)-IGHJ4*01 L123>Q (123) [8.8.21] (1-128)] -trialanil conector (129-131) -[VH humanizado (*Homo sapiens* IGHV3-23*04 (82.50%) -(IGHD)-IGHJ4*01 L123>Q (254) [8.8.21] (132-259)

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EVQLVESGGG LVQPGGSLRL SCAASGRTFS YNPMGWFRQA PGKGRELVAA 50
ISRTGGSTYY PDSVEGRFTI SRDNNAKRMVY LQMNSLRAED TAVYYCAAAG 100
VRAEDGRVRT LPSEYTFWGQ GTQTVVSSAA AEVQLVESGG GLVQPGGSLR 150
LSCAASGRTF SYNPMGWFRQ APGKGRELVA AISRTGGSTY YPDSVEGRFT 200
ISRDNAKRMV YLQMNSLRAE DTAVYYCAA GVRRAEDGRVR TLPSSEYTFWG 250
QGTQVTVSS

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Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-chain 22-96 153-227

cerlapirdinum
cerlapirdine

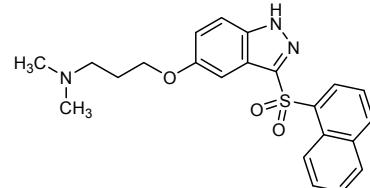
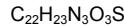
N,N-dimethyl-3-[(3-(naphthalene-1-sulfonyl)-1*H*-indazol-5-yl)oxy]propan-1-amine

cerlapirdine

N,N-diméthyl-3-[(3-(naphthalène-1-sulfonyl)-1*H*-indazol-5-yl)oxy]propan-1-amine

cerlapirdina

N,N-dimetil-3-[(3-(naftaleno-1-sulfonil)-1*H*-indazol-5-il)oxi]propan-1-amina

**dexamcamylaminum**
dexmecamylamine

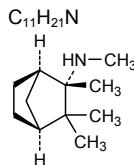
(1*R*,2*S*,4*S*)-*N*,2,3,3-tetramethylbicyclo[2.2.1]heptan-2-amine

dexmécamylamine

(1*R*,2*S*,4*S*)-*N*,2,3,3-tétraméthylbicyclo[2.2.1]heptan-2-amine

dexmecamilamina

(1*R*,2*S*,4*S*)-*N*,2,3,3-tetrametilbicielo[2.2.1]heptan-2-amina

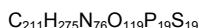


drisapersenum
drisapersen

all-P-ambo-2'-O-methyl-P-thiouridyl-(3'→5')-2'-O-methyl-P-thiocytidyl-(3'→5')-2'-O-methyl-P-thioadenyl-(3'→5')-2'-O-methyl-P-thioguananyl-(3'→5')-2'-O-methyl-P-thioguananyl-(3'→5')-2'-O-methyl-P-thioadenyl-(3'→5')-2'-O-methyl-P-thioguananyl-(3'→5')-2'-O-methyl-P-thioadenyl-(3'→5')-2'-O-methyl-P-thiouridyl-(3'→5')-2'-O-methyl-P-thioadenyl-(3'→5')-2'-O-methyl-P-thioguananyl-(3'→5')-2'-O-methyl-P-thiouridyl-(3'→5')-2'-O-methyl-P-thioguananyl-(3'→5')-2'-O-methyl-P-thiocytidyl-(3'→5')-2'-O-methyl-P-thioadenyl-(3'→5')-2'-O-methyl-P-thiouridyl-(3'→5')-2'-O-methyl-P-thioguananyl-(3'→5')-2'-O-methyl-P-thiocytidyl-(3'→5')-2'-O-methyluridine

drisapersen

drisapersén



(3'-5')-(P-thio)-(Um-Cm-Am-Am-Gm-Gm-Am-Am-Gm-Am-Um-Gm-Gm-Cm-Am-Um-Um-Um-Cm-Um)

faldaprevirum
faldaprevir

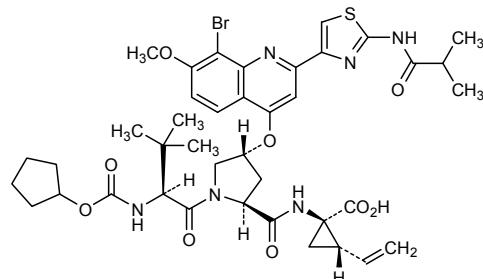
(1*R*,2*S*)-1-{[(2S,4R)-4-[(8-bromo-7-methoxy-2-[2-(methylpropanamido)-1,3-thiazol-4-yl]quinolin-4-yl)oxy]-1-[(2S)-2-[(cyclopentenyl)carbonyl]amino]-3,3-dimethylbutanoyl]pyrrolidine-2-carboxamido}-2-ethenylcyclopropane-1-carboxylic acid

faldaprévir

acide (1*R*,2*S*)-1-[(2*S*,4*R*)-4-[(8-bromo-7-méthoxy-2-[2-(2-méthylpropanamido)-1,3-thiazol-4-yl]quinoléin-4-yl)oxy]-1-[(2*S*)-2-[(cyclopentyloxy)carbonyl]amino]-3,3-diméthylbutanoyl]pyrrolidine-2-carboxamido}-2-éthénylcyclopropane-1-carboxylique

faldaprevir

ácido (1*R*,2*S*)-1-[(2*S*,4*R*)-4-[(8-bromo-7-methoxy-2-[2-(2-metilpropanamido)-1,3-tiazol-4-il]quinolin-4-il)oxi]-1-[(2*S*)-2-[(ciclopentiloxy)carbonil]amino]-3,3-dimetilbutanoil]pirrolidina-2-carboxamido}-2-etenilciclopropano-1-carboxílico

C40H49BrN6O9S

flanvotumabum #

flanvotumab

immunoglobulin G1-kappa, anti-[*Homo sapiens* TYRP1 (tyrosinase-related protein 1, 5,6-dihydroxyindole-2-carboxylic acid oxidase, DHICA oxidase, TRP1, melanoma antigen gp75)], *Homo sapiens* monoclonal antibody; gamma1 heavy chain (1-449) [*Homo sapiens* VH (IGHV7-4-1*02 (95.90%) -(IGHD)-IGHJ4*01) [8.8.12] (1-119) -IGHG1*03 (120-449)], (222-215')-disulfide with kappa light chain (1'-215') [*Homo sapiens* V-KAPPA (IGKV3-11*01 (100.00%) -IGKJ2*01) [6.3.10] (1'-108') -IGKC*01 (109'-215')]; (228-228":231-231")-bisdisulfide dimer

flanvotumab

immunoglobuline G1-kappa, anti-[*Homo sapiens* TYRP1 (protéine 1 apparentée à la tyrosinase, oxydase de l'acide 5,6-dihydroxyindole-2-carboxylique, DHICA-oxydase, TRP1, antigène gp75 du mélanome)], *Homo sapiens* anticorps monoclonal; chaîne lourde gamma1 (1-449) [*Homo sapiens* VH (IGHV7-4-1*02 (95.90%) -(IGHD)-IGHJ4*01) [8.8.12] (1-119) -IGHG1*03 (120-449)], (222-215')-disulfure avec la chaîne légère kappa (1'-215') [*Homo sapiens* V-KAPPA (IGKV3-11*01 (100.00%) -IGKJ2*01) [6.3.10] (1'-108') -IGKC*01 (109'-215')]; dimère (228-228":231-231")-bisdisulfure

flanvotumab

inmunoglobulina G1-kappa, anti-[TYRP1 de *Homo sapiens* (proteína 1 relacionada con la tirosinasa), oxidasa del ácido 5,6-dihidroxiindol-2-carboxílico, DHICA-oxidasa, TRP1, antígeno gp75 de melanoma)], anticuerpo monoclonal de *Homo sapiens*; cadena pesada gamma1 (1-449) [*Homo sapiens* VH (IGHV7-4-1*02 (95.90%) -(IGHD)-IGHJ4*01) [8.8.12] (1-119) -IGHG1*03 (120-449)], (222-215')-disulfuro con la cadena ligera kappa (1'-215') [*Homo sapiens* V-KAPPA (IGKV3-11*01 (100.00%) -IGKJ2*01) [6.3.10] (1'-108') -IGKC*01 (109'-215')]; dímero (228-228":231-231")-bisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada

QVQLVQSGSE LKKPGASVKI SCKASGYTFT SYAMNWRQA PGQGLESMSGW 50
 INTNTGNPTY AQGFTGRFFV SMDTSVSTAY LQISSLKAED TAIYYCAPPY 100
 SSSWYLWDWG QGTLTVVSSA STKGPSVPL APFSKTSGG TAALGCLVKD 150
 YFPEPVTVSW NSGALTSGVH TFPAVLQSSG LYSSLSSVVTW PSSSLGTQTY 200
 ICNVNHKPSN TKVDKRVPEPK SCDKTHTCPP CTAPELLGGP SVFLFPKPK 250
 DTLMISRTPE VTCVVVDVSH EDFEVKENVY VDGVEVHNAK TKPREEQYNS 300
 TYRVSLSLTW LHQDWLNKGK YKCKVSNKAL PAPIEKTISK ARGQPREEQV 350
 YTLPPSREEM TKNQVSLTCL VKGFYPSDIA VEWESENQPE NNYKTPVPL 400
 DSDGSFFLYS KLTVDKSRWQ QGNVFSCCSVN HEALHNHYTQ KSLSLSPGK 449

Light chain / Chaîne légère / Cadena ligera

EIVLTQSPAT LSLSPGERAT LSCRASQSVS SYLAWYQQKP GQAPRLLIYD 50
 ASN RATGIP A RFSGSGSGTD FTLTISLEP EDFAVYVCQQ RSNWLMLYTFG 100
 QGT KLEIKRT VAAPSVPFI FP PSDEQLKSGT ASVUCLNNF YPREAKVQWK 150
 VDNALQSGNS QESVTEQDSK DSTYSLSSTL TLSKADYEKH KVYACEVTHQ 200
 GLSSPVTKSF NRGE 215

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-H 22-96 146-202 263-323 369-427
 22"-96" 146"-202" 263"-323" 369"-427"
 Intra-L 23-88' 135"-195'
 23"-88"" 135"-195"
 Inter-H-L 222-215' 222"-215"
 Inter-H-H 228-228" 231-231"

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
 299, 299"

follitropinum gamma #
follitropin gamma

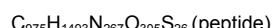
heterodimer of human glycoprotein hormones alpha chain and
 follitropin subunit beta (FSH-beta), follicle stimulating hormone,
 glycoform gamma

follitropine gamma

hétérodimère constitué de la chaîne alpha des hormones
 glycoprotéiques et de la sous-unité bêta de la follitropine (FSH-bêta)
 humaines, hormone folliculostimulante, forme glycosylée gamma

folitropina gamma

heterodímero formado por la cadena alfa de las hormonas
 glicoprotéicas y la subunidad beta de la folitropina (FSH-beta)
 humanas, hormona estimulante del folículo, forma glicosilada
 gamma



Alpha subunit / Sous-unité alpha / Subunidad alfa

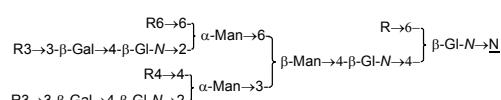
APDVQDCPEC TLQENPFFSQ PGAPILCAG CCFSRAYPTP LRSKKTMLVQ 50
 KNVTSESTCC VAKSYNRVTV MGGFKVENHHT ACHCSTCYYH KS 92

Beta subunit / Sous-unité bêta / Subunidad beta

NSCLENTITI AIEKEECRFC ISINTTWCAG YCYTRDLVYK DPARPKIQKT 50'
 CTFKELVYET VRVPGCAHHA DSYLYTYPVAT QCHCGKCDSD STDCTVRGLG 100'
 PSYCSFGEMK E 111'

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 7-31' 10-60' 28-82' 32-84' 59-87'
 3'-51' 17-66' 20'-104' 28'-82' 32'-84' 87-94'

Glycosylation sites (N) / Sites de glycosylation (N) / Posiciones de glicosilación (N)
 Asn-7' Asn-24' Asn-52 Asn-78



R = α -Fuc or H, R3 = α -Sia or H, R4 and R6 = R3 \rightarrow 3- β -Gal \rightarrow 4- β -GlcNAc or H

gemcitabini elaidas

gemcitabine elaidate

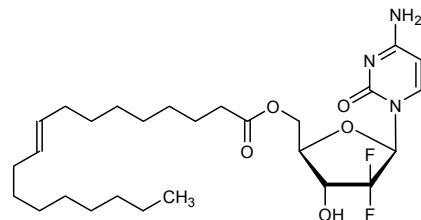
2'-deoxy-2',2'-difluorocytidine 5'-(9E)-octadec-9-enoate

élaïdate de gemcitabine

5'-(9E)-octadéc-9-énoate de 2'-déoxy-2',2'-difluorocytidine

elaídato de gemcitabina

5'-(9E)-octadec-9-enoato de 2'-desoxi-2',2'-difluorocitidina

C₂₇H₄₃F₂N₃O₅**glyceroli phenylbutyras**

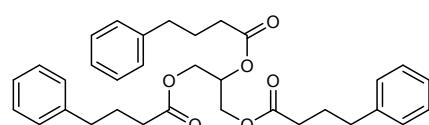
glycerol phenylbutyrate

propane-1,2,3-triyl tris(4-phenylbutanoate)

phénylbutyrate de glycérol

tris(4-phénylbutanoate) de propane-1,2,3-triyle

fenilbutirato de glicerol

C₃₃H₃₈O₆**idursulfasum beta #**

idursulfase beta

iduronate 2-sulfatase (α -L-iduronate sulfate sulfatase), human proenzyme produced in CHO cells (glycoform beta)

idursulfase bêta

iduronate 2-sulfatase (α -L-iduronate sulfate sulfatase), proenzyme humaine produite par des cellules CHO (glycoforme bêta)

idursulfasa beta

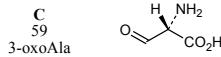
iduronato 2-sulfatasa (α -L-iduronato sulfato sulfatasa), proenzima humana producida por células CHO (forma glicosilada beta)



SETQANSTTD ALNVLLIIVD DLRLPSLGCGC DKLVRSPNID QLASHSLIFQ 50
 NAFAAQAVCA PSRVSFLTGR RPDTTRLYDF NSYWRVHAGN FSTIPQYFKE 100
 NGYVTMSVGK VFHPGISSNH TDDSPYSWSF PPYHPSSSEKY ENTKTCRGPD 150
 GELHANLLCP DVVLVDVPEGT LPDKQSTEQA IQLLEKMKTAS ASPFFLAVGY 200
 HKPHIPFRYP KEFQKLYPLE NITLAPDPEV PGGLPPVAYN PWMDIRQRED 250
 VQALNISVPY GPIPVDFORE IRQSYFASVS YLDTQVGRLL SALDDLQLAN 300
 STIIATFTSDH GWALGEHGEW AKYSNFDVAT HVPLIFYVPG RTASLPEAGE 350
 KLFPLYLDPFD SASQLMEFGR QSMDLVELVS LFFTLAGLAG LQVPPRCPVP 400
 SFHVELCREG KNLLKHFFFR DLEEDPYLPG NFRELIAYSQ YPRPSDIQFW 450
 NSDKPSLKDI KIMCGYSIRTI DYRYTVWVGF NPDEFLANS DIHAGELYVF 500
 DSDPLQDHNM YNDSQGGDLF QLLMP 525

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 146-159 397-407

Modified residue / Résidu modifié / Residuo modificado



Glycosylation sites (N) / Sites de glycosylation (N) / Posiciones de glicosilación (N)
 Asn-6 Asn-90 Asn-119 Asn-221 Asn-255 Asn-300 Asn-488 Asn-512

inlacumab #
inlacumab

immunoglobulin G4-kappa, anti-[*Homo sapiens* SELP (selectin P, CD62)], *Homo sapiens* monoclonal antibody; gamma4 heavy chain (1-451) [*Homo sapiens* VH (IGHV3-13*01 (94.80%) -(IGHD)-IGHJ5*02) [8.7.18] (1-124) -IGHG4*01 hinge S10>P (232), CH2 L1.2>E (239) (125-451)], (138-214')-disulfide with kappa light chain (1'-214') [*Homo sapiens* V-KAPPA (IGKV3-11*01 (100.00%) -IGKJ4*02) [6.3.9] (1'-107') -IGKC*01 (108'-214')]; (230-230':233-233")-bisdisulfide dimer

inlacumab

immunoglobuline G4-kappa, anti-[*Homo sapiens* SELP (sélectine P, CD62)], *Homo sapiens* anticorps monoclonal; chaîne lourde gamma4 (1-451) [*Homo sapiens* VH (IGHV3-13*01 (94.80%) -(IGHD)-IGHJ5*02) [8.7.18] (1-124) -IGHG4*01 charnière S10>P (232), CH2 L1.2>E (239) (125-451)], (138-214')-disulfure avec la chaîne légère kappa (1'-214') [*Homo sapiens* V-KAPPA (IGKV3-11*01 (100.00%) -IGKJ4*02) [6.3.9] (1'-107') -IGKC*01 (108'-214')]; dimère (230-230':233-233")-bisdisulfure

inlacumab

inmunoglobulina G4-kappa, anti-[SELP de *Homo sapiens* (selectina P, CD62)], anticuerpo monoclonal de *Homo sapiens*; cadena pesada gamma4 (1-451) [*Homo sapiens* VH (IGHV3-13*01 (94.80%) -(IGHD)-IGHJ5*02) [8.7.18] (1-124) -IGHG4*01 bisagra S10>P (232), CH2 L1.2>E (239) (125-451)], (138-214')-disulfuro con la cadena ligera kappa (1'-214') [*Homo sapiens* V-KAPPA (IGKV3-11*01 (100.00%) -IGKJ4*02) [6.3.9] (1'-107') -IGKC*01 (108'-214')]; dímero (230-230':233-233")-bisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada
EVQLVESGGG LVRPGGSSLRL SCAASGFTFS NYDMHWVRQA TGKGLEWVSA 50
ITAAAGDIYYP GSVKGRFTIS RENAKNSLYL QMNSLRAGDT AVYYCARGRY 100
SGSGSYNNWD FDPWGQGTIV TVSSASTKGP SVFPLAPCSR STSESTAALG 150
CLVKDYFPEP VTVSNNSGAL TSGVHTFFAV LQSSGLYSLS SVVTVPSSSL 200
GTKTYTCNVD HKPSNTKVDK RVEKYGFPC PPCPAPEFEG GPSVFLFPFK 250
PKDTLMISR PEVTCVVVVDV SQEDPEVQFN WYWDGVVEVHN ARTKPRREEQF 300
NSTYRVVSVL TVLHQDWLNG KEYKCKVSNN GLEFSSTEKT SKAKGQPREP 350
QVYTLPEPQE EMTKNQVSLLT CLVKGFYPSD IAVEWESNGQ PENNYKTTPP 400
VLDSDGSSFL YSRLTVDKSR WQEGNVFSCS VMHEALHNHY TQKSISSLIG 450
K 451

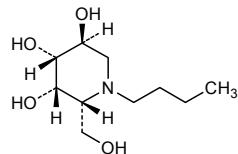
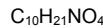
Light chain / Chaîne légère / Cadena ligera
EIVLTQSPAT LSLSFGERAT LSCRASQSVS SYLAWYQQKP GQAPRLLIYD 50
ASN RATGIPA RFSGSGSGTD FTLTISLEP EDFAVYYCQQ RSNWPLTFGG 100
GTKEVIEKRTV AAPSVFIFPP SDEOLKSCTA SVVCLLNNFY PREAKVQWKV 150
DNA LQS GNSQ ESVTEQDSKD STYSLSSTLT LSKADYEKHK VYACEVTHQG 200
LSSPVTKSFN RGE C 214

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
Intra-H 22°-95° 151°-207° 265°-325° 371°-429°
22°-95° 151°-207° 265°-325° 371°-429°
Intra-L 23°-88° 134°-194°
23°-88° 134°-194°
Inter-H-L 138-214° 138°-214°
Inter-H-H 230-230° 233-233°

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
301, 301"

lucerastatum

lucerastat

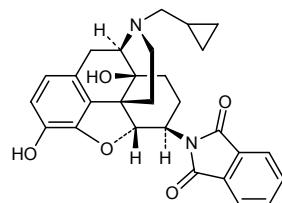
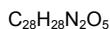
(2*R*,3*S*,4*R*,5*S*)-1-butyl-2-(hydroxymethyl)piperidine-3,4,5-triol**lucérastat**(2*R*,3*S*,4*R*,5*S*)-1-butyl-2-(hydroxyméthyl)pipéridine-3,4,5-triol**lucerastat**(2*R*,3*S*,4*R*,5*S*)-1-butil-2-(hidroximetil)piperidina-3,4,5-triol**natalimidum**

naltalimide

2-[17-(cyclopropylmethyl)-4,5α-epoxy-3,14-dihydroxymorphinan-6β-yl]isoindole-1,3-dione

naltalimide2-[17-(cyclopropylmétihyl)-4,5α-époxy-3,14-dihydroxymorphinan-6β-yl]-2*H*-isoindole-1,3-dione**naltalimida**

2-[17-(ciclopropilmetil)-4,5α-epoxi-3,14-dihidroximorfinan-6β-il]isoindol-1,3-diona



netazepidum
netazepide

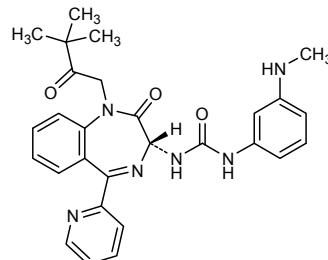
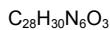
1-[(3*R*)-1-(3,3-dimethyl-2-oxobutyl)-2-oxo-5-(pyridin-2-yl)-2,3-dihydro-1*H*-1,4-benzodiazepin-3-yl]-3-[3-(methylamino)phenyl]urea

nétazépide

1-[(3*R*)-1-(3,3-diméthyl-2-oxobutyl)-2-oxo-5-(pyridin-2-yl)-2,3-dihydro-1*H*-1,4-benzodiazépin-3-yl]-3-[3-(méthylamino)phényl]uree

netazepida

1-[(3*R*)-1-(3,3-dimetil-2-oxobutil)-2-oxo-5-(piridin-2-il)-2,3-dihidro-1*H*-1,4-benzodiazepin-3-il]-3-[3-(metilamino)fenil]urea



niraparibum
niraparib

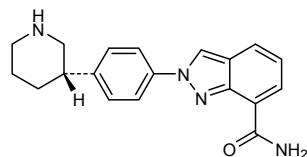
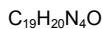
2-{4-[(3*S*)-piperidin-3-yl]phenyl}-2*H*-indazole-7-carboxamide

niraparib

2-{4-[(3*S*)-pipéridin-3-yl]phényl}-2*H*-indazole-7-carboxamide

niraparib

2-{4-[(3*S*)-piperidin-3-il]fenil}-2*H*-indazol-7-carboxamida



ondelopranum

ondelopran

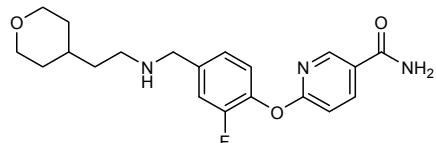
6-[2-fluoro-4-({[2-(oxan-4-yl)ethyl]amino}methyl)phenoxy]pyridine-3-carboxamide

ondélopran

6-[2-fluoro-4-({[2-(oxan-4-yl)éthyl]amino}méthyl)phénoxy]pyridine-3-carboxamide

ondeloprán

6-[2-fluoro-4-({[2-(oxan-4-il)etil]amino}metil)fenoxi]piridina-3-carboxamida

C20H24FN5O3**patiromerum calcium**

patiromer calcium

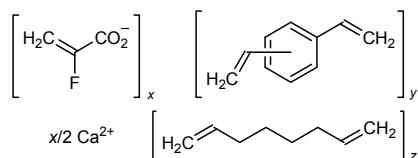
cross-linked polymer of calcium 2-fluoroprop-2-enoate with diethenylbenzene and octa-1,7-diene

patiromère calcique

polymère réticulé de 2-fluoroprop-2-énoate de calcium avec du diéthénylebenzène et de l'octa-1,7-diène

patirómero cálcico

polímero reticulado de 2-fluoroprop-2-enoato de calcio con dietenilbenceno y octa-1,7-dieno

 $[(\text{C}_3\text{H}_2\text{FO}_2)_2 \text{Ca}]_x [\text{C}_8\text{H}_{14}] [\text{C}_{10}\text{H}_{10}]_n$ **patritumab #**

patritumab

immunoglobulin G1-kappa, anti-[*Homo sapiens* ERBB3 (récepteur tyrosine-protéine kinase erbB-3, HER3)], *Homo sapiens* monoclonal antibody;
 gamma1 heavy chain (1-447) [*Homo sapiens* VH (IGHV4-34*01 (99.00%) -(IGHD)-IGHJ2*01) [8.7.11] (1-117) -IGHG1*03 (118-447)], (220-220')-disulfure avec la chaîne légère kappa (1'-220') [*Homo sapiens* V-KAPPA (IGKV4-1*01 (95.00%) -IGKJ1*01) [12.3.9] (1'-113') -IGKC*01 (114'-220')]; (226-226":229-229")-bisdisulfide dimer

patritumab

immunoglobuline G1-kappa, anti-[*Homo sapiens* ERBB3 (récepteur tyrosine-protéine kinase erbB3, HER3)], *Homo sapiens* anticorps monoclonal;
 chaîne lourde gamma1 (1-447) [*Homo sapiens* VH (IGHV4-34*01 (99.00%) -(IGHD)-IGHJ2*01) [8.7.11] (1-117) -IGHG1*03 (118-447)], (220-220')-disulfure avec la chaîne légère kappa (1'-220') [*Homo sapiens* V-KAPPA (IGKV4-1*01 (95.00%) -IGKJ1*01) [12.3.9] (1'-113') -IGKC*01 (114'-220')]; dimère (226-226":229-229")-bisdisulfure

patritumab

inmunoglobulina G1-kappa, anti-[*Homo sapiens* ERBB3 (receptor de tirosina-proteína kinasa erbB3, HER3)], anticuerpo monoclonal de *Homo sapiens*; cadena pesada gamma1 (1-447) [*Homo sapiens* VH (IGHV4-34*01 (99.00%) -(IGHD)-IGHJ2*01) [8.7.11] (1-117) -IGHG1*03 (118-447)], (220-220')-disulfuro con la cadena ligera kappa (1'-220') [*Homo sapiens* V-KAPPA (IGKV4-1*01 (95.00%) -IGK1*01) [12.3.9] (1'-113') -IGKC*01 (114-220')]; dímero (226-226":229-229")-bisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada
 QVQLQQWGAG LLKPSETLSI TCAYVGGSFS GYWSWIRQP PGKGLEWIGE 50
 INHSGSTNYA PSLKSRVTIS VETSKNQFSL KLSSVTAADT AVYYCARDKW 100
 TWYFDLWGRG TLTVVSSAST KGPSPFLAP SSKSTSGGTA ALGCLVKDYF 150
 PEPVTVSWSN GALITSGVHTP PAVLQSSGLY SLSSVTVPS SSLGTQTYIC 200
 NVNHKPSNTK VDKRVEPKSC DKTHTCPFC APELLGGPSV FLFPFPKPKDT 250
 LMISRTEPV TPEVFKFNWYVD GVEVHNATK' PREEQYNSTY 300
 RRVSVLTLH QDWLNKEKEYA CKVSNKALPA PIEKTISKAK GQPREPQVYT 350
 LPPSREEMTK NQVSLTCLVW GFYPSIAVE WESNGQPEENN YKTTTPVILDS 400
 DGSFFLYSKL TVDKSRWQQG NVFSCSVMHE ALHNHYTQKS LSLSPGK 447

Light chain / Chaîne légère / Cadena ligera
 DIEMTQSPLS LAVLGERAT INCRSSQSVL YSSSNRNLYA WYQQNPGQPP 50
 KLLIYWASTR ESGVPDRFSG SGSGTFTLT ISSLQAEDVA VYYCQQYYST 100
 PRTFGQGTKV EIKRTVAAPS VFIFPPSDEQ LKSGTASVVC LLNNFYPREA 150
 KVQNKVDNAL QSGNSQESVT EQDSKDSTYS LSSTTLSKA DYEHKHVYAC 200
 EVTHQGLLSSP VTAKSFNRGEC 220

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-H 22-95 144-200 261-321 367-425
 22"-95" 144"-200" 261"-321" 367"-425"
 Intra-L 23'-94' 140'-200'
 23"-94" 140""-200"
 Inter-H-L 220-220' 220"-220"
 Inter-H-H 226-226" 229-229"

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
 297, 297"

plazomicinum

plazomicin

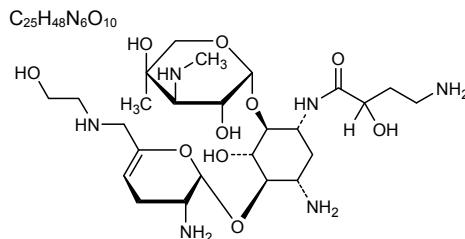
(2S)-4-amino-N-[(1*R*,2*S*,3*S*,4*R*,5*S*)-5-amino-4-[(2*S*,3*R*)-3-amino-6-[(2-hydroxyethyl)amino]methyl]-3,4-dihydro-2*H*-pyran-2-yl]oxy}-2-[(3-deoxy-4-C-methyl-3-(methylamino)-β-L-arabinopyranosyl]oxy}-3-hydroxycyclohexyl]-2-hydroxybutanamide

plazomicine

(2S)-4-amino-N-[(1*R*,2*S*,3*S*,4*R*,5*S*)-5-amino-4-[(2*S*,3*R*)-3-amino-6-[(2-hydroxyéthyl)amino]methyl]-3,4-dihydro-2*H*-pyran-2-yl]oxy}-2-[(3-déoxy-4-C-méthyl-3-(methylamino)-β-L-arabinopyranosyl]oxy}-3-hydroxycyclohexyl]-2-hydroxybutanamide

plazomicina

(2S)-4-amino-N-[(1*R*,2*S*,3*S*,4*R*,5*S*)-5-amino-4-[(2*S*,3*R*)-3-amino-6-[(2-hidroxietil)amino]metil]-3,4-dihidro-2*H*-piran-2-il]oxi}-2-[(3-desoxi-4-C-metil-3-(metilamino)-β-L-arabinopiranosil]oxi}-3-hidroxiciclohexil]-2-hidroxibutanamida



pradigastatum

pradigastat

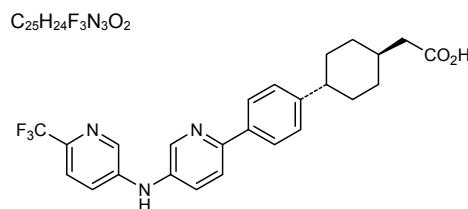
{(1*r*,4*r*)-4-[4-(5-[(6-(trifluoromethyl)pyridin-3-yl]amino)pyridin-2-yl]phenyl)cyclohexyl]acetic acid}

pradigastat

acide *{trans*-4-[4-(5-[6-(trifluorométhyl)pyridin-3-yl]amino)pyridin-2-yl]phényl]cyclohexyl}acétique

pradigastat

ácido {(1r,4r)-4-[4-(5-[(6-(trifluorometil)piridin-3-il)amino]piridin-2-il)fenil]ciclohexil}acético



pritelivirum

pritelivir

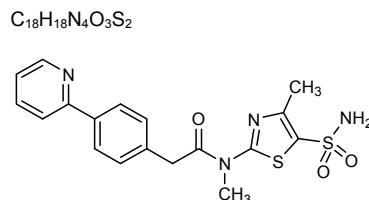
N-methyl-*N*-(4-methyl-5-sulfamoyl-1,3-thiazol-2-yl)-2-[4-(pyridin-2-yl)phenyl]acetamide

pritelivir

N-méthyl-*N*-(4-méthyl-5-sulfamoyl-1,3-thiazol-2-yl)-2-[4-(pyridin-2-yl)phényl]acétamide

pritelivir

N-metil-*N*-(4-metil-5-sulfamoil-1,3-tiazol-2-il)-2-[4-(pyridin-2-il)fenil]acetamida



quilizumabum #

quilizumab

immunoglobulin G1-kappa, anti-[*Homo sapiens* IGHE connecting region (CO) M1 prime (in alternatively spliced heavy chain of membrane IgE on B cells)], humanized monoclonal antibody; gamma1 heavy chain (1-447) [humanized VH (*Homo sapiens*IGHV3-48*01 (85.70%)-IGHD)-IGHJ3*01 M123>L (112) [8.8.10] (1-117) -*Homo sapiens* IGHG1*03 CH1 R120>K (214) (118-447)], (220-219')-disulfide with kappa light chain (1'-219') [humanized V-KAPPA (*Homo sapiens* IGKV1-39*01 (80.00%)-IGKJ1*01) [11.3.9] (1'-112') -*Homo sapiens* IGKC1*01 (113'-219')]; (226-226":229-229")-bisdisulfide dimer.

quilizumab

immunoglobuline G1-kappa, anti-*[Homo sapiens* IgHE région de connexion (CO) M1 prime (dans la chaîne lourde des IgE membranaires à la surface des lymphocytes B, épissée de manière alternative)), anticorps monoclonal humainisé; chaîne lourde gamma1 (1-447) [VH humanisé (*Homo sapiens*IGHV3-48*01 (85.70%) -(IGHD)-IGHJ3*01 M123>L (112) [8.8.10] (1-117) -*Homo sapiens* IGHG1*03 CH1 R120>K (214) (118-447)], (220-219')-disulfure avec la chaîne légère kappa (1'-219') [V-KAPPA humanisé (*Homo sapiens* IGKV1-39*01 (80.00%) -IGKJ1*01) [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113'-219')]; dimère (226-226...229-229")-bisdisulfure

quilizumab

inmunoglobulina G1-kappa, anti-[*Homo sapiens* IGHE región de conexión (CO) M1 prime (en la cadena pesada de las IgE de membrana de la superficie de los linfocitos B, ensamblada de modo alternativo)], anticuerpo monoclonal humanizado; cadena pesada gamma1 (1-447) [VH humanizado (*Homo sapiens*IGHV3-48*01 (85.70%) -(IGHD)-IGHJ3*01 M123>L (112) [8.8.10] (1-117) -*Homo sapiens* IGHG1*03 CH1 R120>K (214) (118-447)], (220-219)-disulfuro con la cadena ligera kappa (1'-219') [V-KAPPA humanizado (*Homo sapiens* IGKV1-39*01 (80.00%) -IGKJ1*01) [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113'-219')]; dímero (226-226"-229-229")-bisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada

Heavy chain	Chain fourc / Cadena pesada
EVQLVSESGGG	LQPGGSSLRL SCASAGFTS
ISDLAYTIIY	ADTVTGRTFI SRDNDSKNTLY
WDAMDWYGCG	TLTVTSSAST GKPGSVFLAP
PEPVTVSWNS	GALTSGVHTF PAVLQSSGLY
VNNHKSPTNK	CDKTHTCPPKC
MLSRSTPEV	VDDVVDSHED PEVKFVNWYD
RVVSVLTVLH	QDWLNGKEYK CKVSNKALPA
LPPSREEMTK	NQVSLLTCLVK GFYPSDIAVE
DGSFFYLSKL	TVDKSRWQQG NIVFCSVVMHE

Light chain / Chaîne légère / Cadena ligera

DIQMTQSPSS	LSASVGRVT	ITCRSSOLV	HNNANTYLHW	YQQKPGKAPK	50
LLIYKVNSRF	SGVPSRSGFS	GSGTDFLTLT	SSLQPEDFAT	YYCSQNTLVP	100
WTFGGGTKVE	IKRTVAAFPSV	FIFPPSDEQL	KGSATSVCL	LNNFYFPREK	150
VTHQGLSSFPV	TGSKNSQEVSTE	QDSKDSTYSL	SSTLTLSKAD	YEKHVKVYACE	200
					219

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-H 22-96 144-200 261-321 367-425
 22-96 " 144-200" 261-321" 367-425"

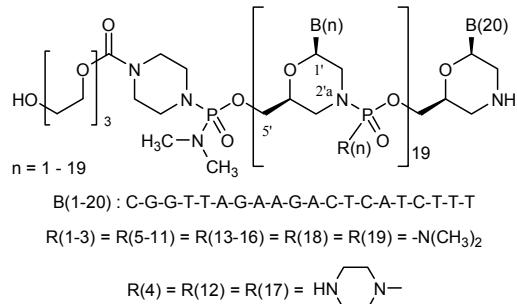
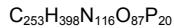
Intra-L 22"-96" 144"-200" 26
 23"-93' 139"-199'
 23""-93"" 139""-199""
 Inter-H-L 220-219" 220"-219"
 Inter-H-H 226-226" 229-229"

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación 297, 297" (non-fucosylated oligosaccharides)

radavirsenum
radavirsen

radavirsen

radavirsén



rafigrelidum
rafigrelide

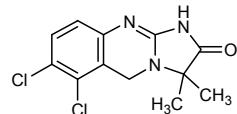
6,7-dichloro-3,3-dimethyl-5,10-dihydroimidazo[2,1-*b*]quinazolin-2(*3H*)-one

rafigrélide

6,7-dichloro-3,3-diméthyl-1,5-dihydroimidazo[2,1-*b*]quinazolin-2(*3H*)-one

rafigrelida

6,7-dicloro-3,3-dimetil-5,10-dihidroimidazo[2,1-*b*]quinazolin-2(*3H*)-ona



refametinibum
refametinib

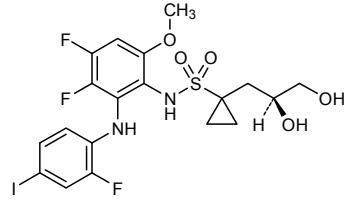
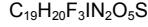
N-(3,4-difluoro-2-[(2-fluoro-4-iodophenyl)amino]-6-methoxyphenyl)-1-[(2*S*)-2,3-dihydroxypropyl]cyclopropane-1-sulfonamide

rémétinib

N-(3,4-difluoro-2-[(2-fluoro-4-iodophényl)amino]-6-méthoxyphényl)-1-[(2*S*)-2,3-dihydroxypropyl]cyclopropane-1-sulfonamide

refametinib

N-(3,4-difluoro-2-[(2-fluoro-4-iodofenil)amino]-6-metoxifenil)-1-[(2*S*)-2,3-dihidroxipropil]ciclopropano-1-sulfonamida



rigosertibum

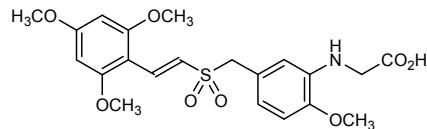
rigosertib

N-[2-methoxy-5-({[(1*E*)-2-(2,4,6-trimethoxyphenyl)ethenyl]sulfonyl}methyl)phenyl]glycine

rigosertib

N-[2-méthoxy-5-({[(1*E*)-2-(2,4,6-triméthoxyphényl)éthenyl]sulfonyl}méthyl)phényl]glycine

rigosertib

N-[2-metoxi-5-({[(1*E*)-2-(2,4,6-trimetoxifenil)etenil]sulfonil}metil)feniil]glicinaC₂₁H₂₅NO₆S**riodipinum**

riodipine

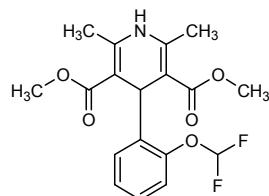
dimethyl 4-[2-(difluoromethoxy)phenyl]-2,6-dimethyl-1,4-dihydropyridine-3,5-dicarboxylate

riodipine

4-[2-(difluorométhoxy)phényl]-2,6-diméthyl-1,4-dihydropyridine-3,5-dicarboxylate de diméthyle

riodipino

4-[2-(difluorometoxi)feniil]-2,6-dimetil-1,4-dihidropiridina-3,5-dicarboxilato de dimetilo

C₁₈H₁₉F₂NO₅**romosozumabum #**

romosozumab

immunoglobulin G2-kappa, anti-[*Homo sapiens* SOST (sclerostin)], humanized monoclonal antibody; gamma2 heavy chain (1-449) [humanized VH (*Homo sapiens* IGHV1-2*02 (87.80%) -(IGHD)-IGHJ2*01 R120>Q (115), L123>T (118) [8.8.16] (1-123) -*Homo sapiens* IGHG2*01 (124-449)], (137-214')-disulfide with kappa light chain (1'-214') [humanized V-KAPPA (*Homo sapiens* IGKV1-33*01 (89.50%) -IGKJ4*02) [6.3.9] (1'-107') -*Homo sapiens* IGKC*01 (108'-214')]; (225-225":226-226":229-229":232-232")-tetrakisdisulfide dimer

romosozumab

immunoglobuline G2-kappa, anti-[*Homo sapiens* SOST (sclérostine)], anticorps monoclonal humanisé; chaîne lourde gamma2 (1-449) [VH humanisé (*Homo sapiens* IGHV1-2*02 (87.80%) -(IGHD)-IGHJ2*01 R120>Q (115), L123>T (118) [8.8.16] (1-123) -*Homo sapiens* IGHG2*01 (124-449)], (137-214')-disulfure avec la chaîne légère kappa (1'-214') [V-KAPPA humanisé (*Homo sapiens* IGKV1-33*01 (89.50%) -IGKJ4*02) [6.3.9] (1'-107') -*Homo sapiens* IGKC*01 (108'-214')]; dimère (225-225":226-226":229-229":232-232")-tétrakisdisulfure

romosozumab

immunoglobulina G2-kappa, anti-[SOST (esclerostina) de *Homo sapiens*], anticuerpo monoclonal humanizado; cadena pesada gamma2 (1-449) [VH humanizado (*Homo sapiens* IGHV1-2*02 (87.80%) -(IGHD)-IGHJ2*01 R120>Q (115), L123>T (118) [8.8.16] (1-123) -*Homo sapiens*IGHG2*01 (124-449)], (137-214')-disulfuro con la cadena ligera kappa (1'-214') [V-KAPPA humanizado (*Homo sapiens* IGKV1-33*01 (89.50%) -IGKJ4*02) [6.3.9] (1-107') -*Homo sapiens*IGKC*01 (108'-214')]; dímero (225-225":226-226":229-229":232-232")-tetrakisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada
EVQLVQSGAE VKKEPGASVKV SCKASGYTFT DYNMHWVRQA PGQGLEWME 50
INPNNSGGAGY NQKFKGRVTM TTDTSSTSTAY MELRSLRSDD TAVYYCARLG 100
YDDIYDWWYF DVWGQQGTTVT VSSASTKGPS VFVFLAPCSRS TSESTAALGC 150
LVKDYFPEPV TTVSNNSGALT SGVHTTFAVL QSSGLYSLSV VVTVEFSNFG 200
TQTYTCNVDH KPSNTKVDKT VERKCCFVCP PCPAPPVAGP SVFLFPPPK 250
DTLMISRPTPE VTCVVVDVDSH EDEPEVQFNWY VDGVEVHNAK TKPREEQFNS 300
TFRVVSLLTV VHQDWLNKGK YKCKVSNKGL PAPIEKTIISK TKGQPREPQV 350
YTLPPSREEM TKNQVSLTCL VKGFYPSDIA VEWESNGQPE NNYKTTPPML 400
DSDGSFFLYS KLTVDKSRWQ QGNVFSCSVN HEALHNHYTQ KSLSLSPGK 449

Light chain / Chaîne légère / Cadena ligera
DIQMKTQSPSS LSASVGDRTV ITCRASQDIS NYLNWYQQKP GKAPKLLIYY 50
TSRLLLSGVPS RFSGSSGGTD FTLTISLQP EDFATYYCQQ GDTLPYTFGG 100
GTVKEIKRTV AAPSVFIFPP SDEQLIKSGTA SVVCLLNNFY FREAKVQWKV 150
DNAQSGNSQ ESVTEQDSKD STYSLSSLT LSKADYEKKH VYACEVTHQG 200
LSSPVTKSFN RGEc 214

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
Intra-H 22-96 150-206 263-323 369-427
22"-96" 150"-206" 263"-323" 369"-427"
Intra-L 23-88 134"-194'
23"-88" 134"-194"
Inter-H-L 137-214' 137"-214"
Inter-H-H 225-225" 226-226" 229-229" 232-232"

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
299, 299"

samidorphanum
samidorphan

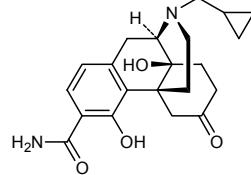
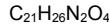
17-(cyclopropylmethyl)-4,14-dihydroxy-6-oxomorphinan-3-carboxamide

samidorphan

17-(cyclopropylmethyl)-4,14-dihydroxy-6-oxomorphinan-3-carboxamide

samidorfano

17-(ciclopropilmetil)-4,14-dihidroxi-6-oxomorfinan-3-carboxamida

**sapitinibum**
sapitinib

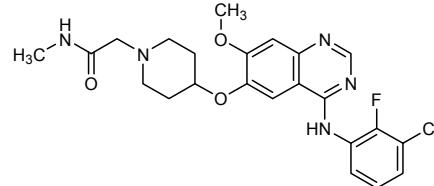
2-[4-({4-[(3-chloro-2-fluorophenyl)amino]-7-methoxyquinazolin-6-yl}oxy)piperidin-1-yl]-N-methylacetamide

sapitinib

2-[4-({4-[(3-chloro-2-fluorophényle)amino]-7-méthoxyquinazolin-6-yl}oxy)pipéridin-1-yl]-N-méthylacétamide

sapitinib

2-[4-({4-[(3-cloro-2-fluorofenil)amino]-7-metoxiquinazolin-6-il}oxi)piperidin-1-il]-N-metilacetamida

**sarilumab #**
sarilumab

immunoglobulin G1-kappa, anti-[*Homo sapiens* IL6R (interleukin receptor, IL-6R, CD126)], *Homo sapiens* monoclonal antibody; gamma1 heavy chain (1-446) [*Homo sapiens* VH (IGHV3-9*01 (94.90%) -(IGHD)-IGHJ3*02) [8.8.9] (1-116) -IGHG1*01 (117-446)], (219-214')-disulfide with kappa light chain (1'-214') [*Homo sapiens* V-KAPPA (IGKV1-12*01 (96.80%) -IGKJ2*01) [6.3.9] (1'-107') -IGKC*01 (108'-214')]; (225-225":228-228")-bisdisulfide dimer

sarilumab

immunoglobuline G1-kappa, anti-[*Homo sapiens* IL6R (récepteur de l'interleukine 6, IL-6R, CD126)], *Homo sapiens* anticorps monoclonal; chaîne lourde gamma1 (1-446) [*Homo sapiens* VH (IGHV3-9*01 (94.90%) -(IGHD)-IGHJ3*02) [8.8.9] (1-116) -IGHG1*01 (117-446)], (219-214')-disulfure avec la chaîne légère kappa (1'-214') [*Homo sapiens* V-KAPPA (IGKV1-12*01 (96.80%) -IGKJ2*01) [6.3.9] (1'-107') -IGKC*01 (108'-214')]; dimère (225-225":228-228")-bisdisulfure

sarilumab

inmunoglobulina G1-kappa, anti-[*Homo sapiens* IL6R (receptor de la interleukina 6, IL-6R, CD126)], anticuerpo monoclonal de *Homo sapiens*; cadena pesada gamma1 (1-446) [*Homo sapiens* VH (IGHV3-9*01 (94.90%) -(IGHD)-IGHJ3*02) [8.8.9] (1-116) -IGHG1*01 (117-446)], (219-214')-disulfuro con la cadena ligera kappa (1'-214') [*Homo sapiens* V-KAPPA (IGKV1-12*01 (96.80%) -IGKJ2*01) [6.3.9] (1'-107') -IGKC*01 (108'-214')]; dímero (225-225":228-228")-bisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada

EVQLVESGGG LVQPGRLSLR SCAASRFTFD DYAMHWVRQA PGKGLEWVSG 50
 ISWNNSGRIGY ADSVKGRFTI SRDNAEENSLF LQMNGLRAED TALYYCAKGR 100
 DSFDIWGQCT MVTVSSASTK GPSVFPLAPS SKSTSGGTAA LGCLVKDYFP 150
 EPVTVSWNSG ALTSGVHTFP AVLQSSGLYS LSSVVTVFSS SLGTQTYICN 200
 VNHKPSNTKRV DKKVEPKSCD KTHTCPCPFA PELLGGPSVLF 250
 MISRTPEVITC VVVDVSHEDP EVKFNWVYVDG VEVHNAKTK REEQYNSTYR 300
 VVSVLTVLHQ DWLNKEYKC KVSNKALPAP IEKTISKAKG QPREPVQYTL 350
 PPSRDELTKN QVSLTCLVKG FYPSDIAEW ESNGOPENNY KTTPPVLDSD 400
 GSFFLYSKLT VDKSRWQQGN VFSCSVMHEA LHNHYTQKSL SLSPGK 446

Light chain / Chaîne légère / Cadena ligera

DIGMTQSPSS VSAVGRDVT ITCRASQGIS SWLAWYQQKP GKAPKLLIYG 50
 ASSLESGVPS RFSGSGSGTD FTILTISSLQP EDFASYYCQQ ANSFPTYTFQ 100
 GTKLEIKRTV AAPSVFIFPP SDEQLKSGTA SVVCLLNFFY PREAKVQWKV 150
 DNALQSGNSQ ESVTEQDSKD STYSLSSTLT LSKADYEKHK VYACEVTHQG 200
 LSSPVTKSFn RGEc 214

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro

Intra-H 22-96 143-199 260-320 366-424

22"-96" 143"-199" 260"-320" 366"-424"

Intra-L 23"-88' 134"-194'

23"-88" 134"-194"

Inter-H-L 219-214' 219"-214"

Inter-H-H 225-225" 228-228"

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación

296, 296"

secretinum humanum
secretin human

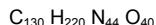
human peptide hormone secretin;
 L-histidyl-L-seryl-L-aspartylglycyl-L-threonyl-L-phenylalanyl-L-threonyl-L-seryl-L-glutamyl-L-leucyl-L-seryl-L-arginyl-L-leucyl-L-arginyl-L-glutamylglycyl-L-alanyl-Larginyl-L-leucyl-L-glutaminy-L-arginyl-L-leucyl-L-leucyl-L-glutaminylglycyl-L-leucyl-L-valinamide

sécrétine humaine

sécrétine humaine hormone peptidique;
 L-histidyl-L-séryl-L-aspartylglycyl-L-thréonyl-L-phénylalanyl-L-thréonyl-L-séryl-L-glutamyl-L-leucyl-L-séryl-L-arginyl-L-leucyl-L-arginyl-L-glutamylglycyl-L-alanyl-L-arginyl-L-leucyl-L-glutaminy-L-arginyl-L-leucyl-L-leucyl-L-glutaminylglycyl-L-leucyl-L-valinamide

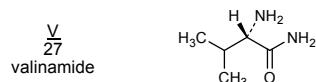
secretina humana

secretina humana, hormona peptídica;
 L-histidil-L-seril-L-aspartilglicil-L-treonil-L-fenilalanil-L-treonil-L-seril-L-glutamil-L-leucil-L-seril-L-arginil-L-leucil-L-arginil-L-glutamiglicil-L-alanil-L-arginil-L-leucil-L-glutaminil-L-arginil-L-leucil-L-leucil-L-glutaminilglicil-L-leucil-L-valinamida



HSDGTFSEL SRLREGARLQ RLLQGLV 27

Modified residue / Résidu modifié / Residuo modificado

**selisistatum**
selisistat

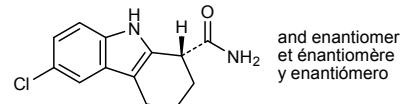
rac-6-chloro-2,3,4,9-tetrahydro-1*H*-carbazole-1-carboxamide

sélisistat

rac-6-chloro-2,3,4,9-tétrahydro-1*H*-carbazole-1-carboxamide

selisistat

rac-6-cloro-2,3,4,9-tetrahidro-1*H*-carbazol-1-carboxamida

**setrobuvirum**
setrobuvir

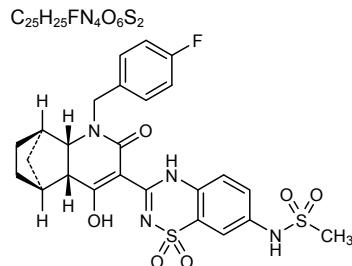
N-(3-[(4*aR*,5*S*,8*R*,8*aS*)-1-[(4-fluorophenyl)methyl]-4-hydroxy-2-oxo-1,2,4*a*,5,6,7,8,8*a*-octahydro-5,8-methanoquinolin-3-yl]-1,1-dioxo-1,4-dihydro-1*λ*⁶,2,4-benzothiadiazin-7-yl)methanesulfonamide

sétrobuvir

N-(3-[(4*aR*,5*S*,8*R*,8*aS*)-1-[(4-fluorophényl)méthyl]-4-hydroxy-2-oxo-1,2,4*a*,5,6,7,8,8*a*-octahydro-5,8-méthanoquinoléin-3-yl]-1,1-dioxo-1,4-dihydro-1*λ*⁶,2,4-benzothiadiazin-7-yl)méthanesulfonamide

setrobuvir

N-(3-[(4*aR*,5*S*,8*R*,8*aS*)-1-[(4-fluorofenil)metil]-4-hidroxi-2-oxo-1,2,4*a*,5,6,7,8,8*a*-octahidro-5,8-metanoquinolin-3-il]-1,1-dioxo-1,4-dihidro-1*λ*⁶,2,4-benzotiadiazin-7-il)metanosulfonamida



sevuparinum natricum
sevuparin sodium

sodium salt of a low molecular mass heparin obtained by depolymerization through periodate oxidation of heparin from porcine intestinal mucosa, followed by reduction and mild acid hydrolysis of the product; the majority of the components have a 2-amino-2-deoxy-D-glucopyranose derivative structure at both ends of their chain, the one at the reducing end can be substituted with threonic acid or erythronic acid; the relative average molecular mass range is approximately 7,500 daltons with about 90% ranging between 2,000 and 15,000 daltons; the degree of sulfation is 2 to 2.5 per disaccharide unit

sévuparine sodique

sel sodique d'une héparine de basse masse moléculaire obtenue par dépolymérisation, au moyen d'une oxydation périodique, d'héparine de muqueuse intestinale de porc, suivi par une réduction et par une hydrolyse douce; la majorité des composants de la sévuparine sodique possèdent une structure 2-amino-2-déoxy-D-glucopyranose aux deux extrémités, la réductrice peut être substituée par un acide thrénique ou érythronique; la masse moléculaire relative moyenne est approximativement de 7500 daltons, et celles de 90% sont comprises entre 2000 à 15000; le degré de sulfatation est de 2 à 2,5 par unité disaccharide.

sevuparina sódica

sal sódica de una heparina de baja masa molecular obtenida por despolimerización, mediante oxidación periódica, de heparina de mucosa intestinal porcina, seguida de reducción e hidrólisis con ácido débil; la mayoría de los componentes de la sevuparina sódica tienen una estructura 2-amino-2-desoxi-D-glucopiranosa en los dos extremos, el reductor puede estar substituido con un ácido treónico o eritrónico; la masa molecular relativa media es aproximadamente de 7500 dalton, y con el 90% comprendido entre 2000 y 15000; el grado de sulfatación es de 2 a 2,5 por unidad de disacárido

solitomabum #
solitomab

immunoglobulin scFv-scFv, anti-[*Homo sapiens* EPCAM (epithelial cell adhesion molecule, tumor-associated calcium signal transducer 1, TACSTD1, gastrointestinal tumor-associated protein 2, GA733-2, epithelial glycoprotein 2, EGFR, KSA, KS1/4 antigen, M4S1, tumor antigen 17-1A, Ep-CAM, EpCAM, CD326)]/anti-[*Homo sapiens* CD3E (CD3 epsilon)], *Mus musculus* monoclonal antibody bispecific single chain;
scFv anti-EPCAM [*Mus musculus* V-KAPPA (IGKV8-19*01 (98.00%)-IGKJ5*01 L126>I (112)) [12.3.9] (1-113)-15-mer tris(tetraglycyl-seryl) linker (114-128) -*Mus musculus* VH (IGHV1-54*01 (85.90%)-(IGHD)-IGHJ4*01, S123>T (243)) [8.8.14] (129-248)] -5-mer tetraglycyl-seryl linker (249-253) -scFv anti-CD3E [humanized VH (*Homo sapiens* IGHV1-46*01 (82.50%)-(IGHD)-IGHJ6*01) [8.8.12] (254-372) -18-mer linker (373-390) -V-KAPPA (*Mus musculus* IGKV4-59*01 (81.70%)-IGKJ1*01 L124>V (493) [5.3.9] (391-496)] -hexahistidine (497-502)

solitomab	immunoglobuline scFv-scFv, anti-[<i>Homo sapiens</i> EPCAM (molécule d'adhésion des cellules épithéliales, transducteur 1 du signal calcium associé aux tumeurs, TACSTD1, protéine 2 associée aux tumeurs gastrointestinales, GA733-2, glycoprotéine épithéliale 2, EGP-2, KSA, antigène KS1/4, M4S1, antigène tumoral 17-1A, Ep-CAM, EpCAM, CD326)]/anti-[<i>Homo sapiens</i> CD3E (CD3 epsilon)], <i>Mus musculus</i> anticorps monoclonal biespécifique à chaîne unique; scFv anti-EPCAM [<i>Mus musculus</i> V-KAPPA (IGKV8-19*01 (98.00%)-IGKJ5*01 L126>I (112)) [12.3.9] (1-113) -15-mer tris(tétraglycyl-séryl) linker (114-128) - <i>Mus musculus</i> VH (IGHV1-54*01 (85.90%)-(IGHD)-IGHJ4*01, S123>T (243)) [8.8.14] (129-248)] -5-mer tétraglycyl-séryl linker (249-253) -scFv anti-CD3E [VH humanisé (<i>Homo sapiens</i> IGHV1-46*01 (82.50%)-(IGHD)-IGHJ6*01) [8.8.12] (254-372) -18-mer linker (373-390) -V-KAPPA (<i>Mus musculus</i> IGKV4-59*01 (81.70%)-IGKJ1*01 L124>V (493) [5.3.9] (391-496)] -hexahistidine (497-502)
solitomab	inmunoglobulina scFv-scFv, anti-[EPCAM de <i>Homo sapiens</i> (molécula de adhesión de células epiteliales, transductor 1 de la señal de calcio asociado a tumores, TACSTD1, proteína 2 asociada a tumores gastrointestinales, GA733-2, glicoproteína epitelial 2, EGP-2, KSA, antigén KS1/4, M4S1, antigén tumoral 17-1A, Ep-CAM, EpCAM, CD326)]/anti-[<i>Homo sapiens</i> CD3E (CD3 epsilon)], anticuerpo monoclonal biespecífico monocatenario de <i>Mus musculus</i> ; scFv anti-EPCAM [<i>Mus musculus</i> V-KAPPA (IGKV8-19*01 (98.00%)-IGKJ5*01 L126>I (112)) [12.3.9] (1-113) -15-mer tris(tetraglicil-séryl) conector (114-128) - <i>Mus musculus</i> VH (IGHV1-54*01 (85.90%)-(IGHD)-IGHJ4*01, S123>T (243)) [8.8.14] (129-248)] -5-mer tetraglicil-séryl conector (249-253) -scFv anti-CD3E [VH humanizado (<i>Homo sapiens</i> IGHV1-46*01 (82.50%)-(IGHD)-IGHJ6*01) [8.8.12] (254-372) -18-mer conector (373-390) -V-KAPPA (<i>Mus musculus</i> IGKV4-59*01 (81.70%)-IGKJ1*01 L124>V (493) [5.3.9] (391-496)] -hexahistidina (497-502)

ELVMTQSPSS LTVTAGEKVT MSCKSSQSLL NSGNQKNYLT WYQQKPGQPP 50
 KLLIQWASTR ESGVPDRFTG SGSGTDFTL ISSVQAEGLA VYYCQNDSY 100
 PLTFAGTTLK EIKGGGGSGG GGSGGGGSEV QLLEOSGAEL VRPGTSVKIS 150
 CKASGYAFTN YWLGWVKQRP GHGLEWIGDI FPFGSGNIHYN EKFKGKAILT 200
 ADKSSSTAYM QLSSLTFEDS AVYFCARLRN WDEPMDYWGQ GTTVTVSSGG 250
 GGSQVQLVQS GAEVKKPGAA VKVSCKASGY TFTRYTMHWV RQAPGQGLEW 300
 IGYINPSRGY TNYADSVKGR FTITTDKSTS TAYMELSSLR SEDTATYYCA 350
 RYYDDHYCLD YWGGOTTVTV SSGEGTSTGS GGSGGGGAG DIVLTQSPAT 400
 LSLSPERAT LSCRASQSVS YMNWYQKPG KAPKRWIYDT SKVASCPAR 450
 FSGSGSGTGY SLTINSLEAE DAATYYCQW SSNPLTFFGG TKVEIKHHH 500
 HH 502

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-chain 23-94 151-225 275-349 413-477

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
 305 (but Pro in 306)

sovaprevirum sovaprevir

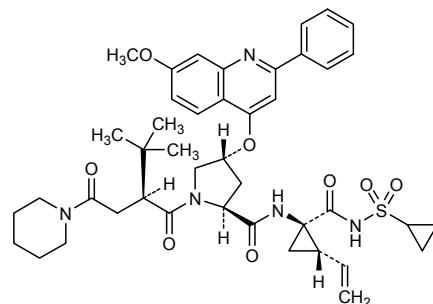
(2*S*,4*R*)-1-[(2*S*)-2-*tert*-butyl-4-oxo-4-(piperidin-1-yl)butanoyl]-
N-{(1*R*,2*S*)-1-[(cyclopropanesulfonyl)carbamoyl]-
 2-ethenylcyclopropyl}-4-[(7-methoxy-2-phenylquinolin-
 4-yl)oxyl]pyrrolidine-2-carboxamide

sovaprévir

(*2S,4R*)-1-[(*2S*)-2-*tert*-butyl-4-oxo-4-(piperidin-1-yl)butanoyl]-
N-(*1R,2S*)-1-[(cyclopropanesulfonyl)carbamoyl]-
 2-éthyénylcyclopropyl]-4-[(7-méthoxy-2-phénylquinoléin-4-yl)oxy]pyrrolidine-2-carboxamide

sovaprevir

(*2S,4R*)-1-[(*2S*)-2-*tert*-butyl-4-oxo-4-(piperidin-1-yl)butanoyl]-
N-(*1R,2S*)-1-[(cyclopropanesulfonyl)carbamoyl]-2-etenilciclopropil]-4-[(7-metoxi-2-fenilquinolin-4-yl)oxy]pirrolidina-2-carboxamida

C43H53N5O8S**sutezolidum**

sutezolid

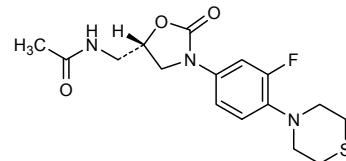
N-{(*5S*)-3-[3-fluoro-4-(thiomorpholin-4-yl)phenyl]-2-oxo-1,3-oxazolan-5-yl}methyl)acetamide

sutézolid

N-{(*5S*)-3-[3-fluoro-4-(thiomorpholin-4-yl)phényl]-2-oxo-1,3-oxazolidin-5-yl}méthyl)acétamide

sutezolid

N-{(*5S*)-3-[3-fluoro-4-(tiomorfolin-4-yl)fenil]-2-oxo-1,3-oxazolan-5-yl}metil)acetamida

C16H20FN3O3S**tanzisertibum**

tanzisertib

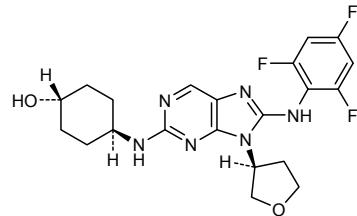
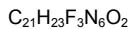
(*1r,4r*)-4-{9-[(*3S*)-oxolan-3-yl]-8-[(2,4,6-trifluorophenyl)amino]-9*H*-purin-2-yl}amino)cyclohexan-1-ol

tanzisertib

(*1r,4r*)-4-{9-[(*3S*)-oxolan-3-yl]-8-[(2,4,6-trifluorophényl)amino]-9*H*-purin-2-yl}amino)cyclohexan-1-ol

tanzisertib

(*1r,4r*)-4-{9-[(*3S*)-oxolan-3-yl]-8-[(2,4,6-trifluorfenil)amino]-9*H*-purin-2-yl}amino)cyclohexan-1-ol



tavaborolum
tavaborole

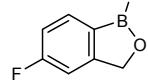
5-fluoro-2,1-benzoxaborol-1(3*H*)-ol

tavaborole

5-fluoro-2,1-benzoxaborol-1(3*H*)-ol

tavaborol

5-fluoro-2,1-benzoxaborol-1(3*H*)-ol



tedatioxetinum
tedatioxetine

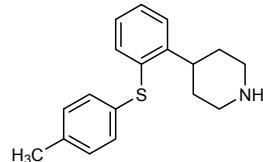
4-{2-[{(4-methylphenyl)sulfanyl]phenyl}piperidine

tédatioxétine

4-{2-[{(4-méthylphényle)sulfanyl]phényle}pipéridine

tedatioxetina

4-{2-[{(4-metilfenil)sulfanil]fenil}piperidina



tipiracilum
tipiracil

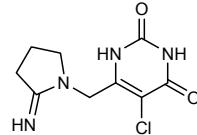
5-chloro-6-[(2-iminopyrrolidin-1-yl)methyl]pyrimidine-2,4-(1*H*,3*H*)-dione

tipiracil

5-chloro-6-[(2-iminopyrrolidin-1-yl)méthyl]pyrimidine-2,4-(1*H*,3*H*)-dione

tipiracilo

5-cloro-6-[(2-iminopirrolidin-1-il)metil]pirimidina-2,4-(1*H*,3*H*)-diona



tirasemtivum

tirasemtiv

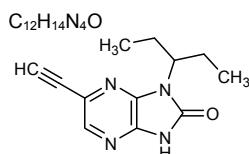
6-ethynyl-1-(pentan-3-yl)-2H-imidazo[4,5-b]pyrazin-2-one

tirasemtiv

6-éthynyl-1-(pentan-3-yl)-1,3-dihydro-2H-imidazo[4,5-b]pyrazin-2-one

tirasemtiv

6-etinil-1-(pentan-3-il)-2H-imidazo[4,5-b]pirazin-2-oná

**tozadenantum**

tozadenant

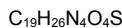
4-hydroxy-N-[4-methoxy-7-(morpholin-4-yl)-1,3-benzothiazol-2-yl]-4-methylpiperidine-1-carboxamide

tozadénant

4-hydroxy-N-[4-méthoxy-7-(morpholin-4-yl)-1,3-benzothiazol-2-yl]-4-méthylpipéridine-1-carboxamide

tozadenant

4-hidroxi-N-[4-metoxi-7-(morfolin-4-il)-1,3-benzotiazol-2-il]-4-metilpiperidina-1-carboxamida

**trebananibum #**

trebananib

immunoglobulin G1 Fc fragment fused with two synthetic polypeptides that bind the *Homo sapiens* ANGPT2 (angiopoietin 2); methionyl (1)-gamma1 heavy chain fragment (2-228) [*Homo sapiens* IGHG1*01 hinge (EPKSC 1-5>del) (2-11), CH2 (12-121), CH3 (122-228)] fused, at the C-terminal end, with a synthetic polypeptide that comprises two 14-mer amino acid repeats that bind angiopoietin 2 (229-287) [linker (229-235) -14-mer (236-249) -linker (250-271) -14-mer (272-285) -leucyl-glutamate]; (7-7':10-10')-bisdisulfide dimer

trébananib

immunoglobuline G1 fragment Fc fusionné à deux polypeptides synthétiques qui se lient à l'*Homo sapiens* ANGPT2 (angiopoïétine 2); méthionyl (1)-fragment de chaîne gamma1 (2-228) [*Homo sapiens* IGHG1*01 charnière (EPKSC 1-5>del) (2-11), CH2 (12-121), CH3 (122-228)] fusionné, à l'extrémité C-terminale, à un polypeptide synthétique qui comprend deux motifs répétés de 14 acides aminés qui se lient à l'angiopoïétine 2 (229-287) [linker (229-235) -14-mer (236-249) -linker (250-271) -14-mer (272-285) -leucyl-glutamate]; dimère (7-7':10-10')-bisdisulfure

trebananib

inmunoglobulina G1 fragmento Fc fusionado con dos polipéptidos sintéticos que se unen a la ANGPT2 (angiopoietina 2) de *Homo sapiens*; metionil (1)-fragmento de cadena gamma1 (2-228) [*Homo sapiens* IGHG1*01 bisagra (EPKSC 1-5>del) (2-11), CH2 (12-121), CH3 (122-228)] fusionada con el extremo C-terminal de un polipéptido sintético que comprende dos secuencias repetidas de 14 aminoácidos que se unen a la angiopoietina 2 (229-287) [conector (229-235)-14-mer (236-249)-conector (250-271)-14-mer (272-285)-leucil-glutamato]; dímero (7'-7':10-10')-bisdisulfuro

```

MDKTHTCPPC PAPELLGGPS VFLFPPPKPD TLMISRTPEV TCVVVDSHE 50
DPEVKFNWYV DGVEVHNAKT KPRREEQYNST YRVVSVLTVL HQDWLNGKEY 100
KCKVSNKALP APIEKTISKA KGQPREPOVY TLEPSRDELK KNQVSLTCLV 150
KGFYPSDIAV EWESNGOPEN NYKTTPPVLD SDGSFFLYSK LTVDKSRWQQ 200
GNVFSCVMH EALHNHYTQK SLSLSPGKGG GGGAAQEECE WDPWTCEHMG 250
SGSATGGSGS TASSGSGSAT HQEECEWDWP TCEHMLE 287

```

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-chain 42-102 148-206 239-246 275-282
 42'-102' 148'-206' 239'-246' 275'-282'
 Inter-chains 7-7' 10-10'

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
 78, 78'

trelagliptinum
 treagliptin

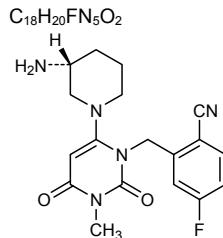
2-({6-[(3*R*)-3-aminopiperidin-1-yl]-3-methyl-2,4-dioxo-3,4-dihydropyrimidin-1(2*H*)-yl)methyl)-4-fluorobenzonitrile

trélagliptine

2-({6-[(3*R*)-3-aminopiperidin-1-yl]-3-méthyl-2,4-dioxo-3,4-dihydropyrimidin-1(2*H*)-yl)méthyl)-4-fluorobenzonitrile

trelagliptina

2-({6-[(3*R*)-3-aminopiperidin-1-yl]-3-metil-2,4-dioxo-3,4-dihidropirimidin-1(2*H*)-yl)metil)-4-fluorobenzonitrilo


umeclidinium bromidum
 umeclidinium bromide

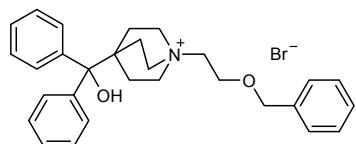
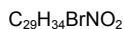
1-{2-[(benzyl)oxy]ethyl}-4-[hydroxydi(phenyl)methyl]-1-azabicyclo[2.2.2]octan-1-ium bromide

bromure d'uméclidinium

bromure de 1-{2-[(benzyl)oxy]éthyl}-4-[hydroxydi(phényl)méthyl]-1-azabicyclo[2.2.2]octanium

bromuro de umeclidinio

bromuro de 1-{2-[(benzil)oxi]etil}4-[hidroxidi(fenil)metil]-1-azabiciclo[2.2.2]octan-1-io

**vapendavirum**

vapendavir

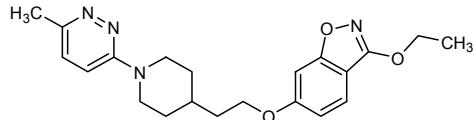
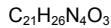
3-ethoxy-6-{2-[1-(6-methylpyridazin-3-yl)piperidin-4-yl]ethoxy}-1,2-benzoxazole

vapendavir

3-éthoxy-6-{2-[1-(6-méthylpyridazin-3-yl)pipéridin-4-yl]éthoxy}-1,2-benzoxazole

vapendavir

3-etoxy-6-{2-[1-(6-metilpiridazin-3-il)piperidin-4-il]etoxi}-1,2-benzoxazol

**vonoprazanum**

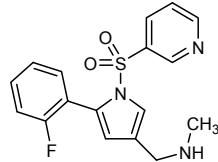
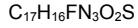
vonoprazan

1-[5-(2-fluorophenyl)-1-(pyridine-3-sulfonyl)-1*H*-pyrrol-3-yl]-*N*-methylmethanamine

vonoprazan

1-[5-(2-fluorophényl)-1-(pyridine-3-sulfonyl)-1*H*-pyrrol-3-yl]-*N*-méthylméthanamine

vonoprazán

1-[5-(2-fluorofenil)-1-(piridina-3-sulfonil)-1*H*-pirrol-3-il]-*N*-metilmelanamina**vortioxetinum**

vortioxetine

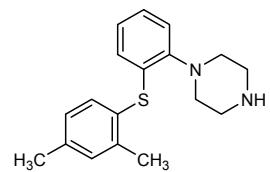
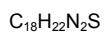
1-{2-[(2,4-dimethylphenyl)sulfanyl]phenyl}piperazine

vortioxétine

1-{2-[(2,4-diméthylphényl)sulfanyl]phényl}pipérazine

vortioxetina

1-{2-[(2,4-dimetilfenil)sulfani]fenil}piperazina

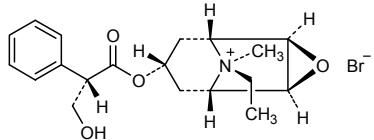


AMENDMENTS TO PREVIOUS LISTS
MODIFICATIONS APPORTÉES AUX LISTES ANTÉRIEURES
MODIFICACIONES A LAS LISTAS ANTERIORES

Recommended International Non Proprietary Names (Rec. INN): List 17
(Chronicle of the WHO, 1977, Vol. 31, No. 10)

p. 7 **oxitropii bromidum**
 oxitropium bromide replace the description and the structure by the following ones

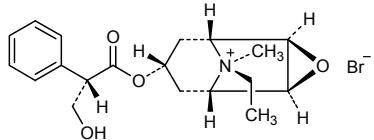
(*1R,2R,4S,5S,7s,9s*)-9-ethyl-7-[(*2S*)-3-hydroxy-2-phenylpropanoyl]oxy}-9-methyl-3-oxa-9-azoniatricyclo[3.3.1.0^{2,4}]nonane bromide



Dénominations communes internationales recommandées (DCI Rec.): Liste 17
(Chronique de l'OMS, Vol. 31, No. 10, 1977)

p. 7 **oxitropii bromidum**
 bromure d'oxitropium remplacer la description et la structure par les suivantes

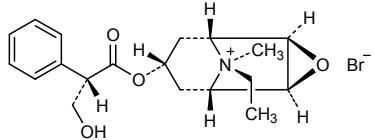
bromure de (*1R,2R,4S,5S,7s,9s*)-9-éthyl-7-[(*2S*)-3-hydroxy-2-phénylpropanoyl]oxy}-9-méthyl-3-oxa-9-azoniatricyclo[3.3.1.0^{2,4}]nonane



Denominaciones Comunes Internacionales Recomendadas (DCI Rec.): Lista 17
(Crónica de la OMS, Vol. 31, No. 10, 1977)

p. 7 **oxitropii bromidum**
 bromuro de oxitropio sustitúyase la descripción y la estructura por las siguientes

bromuro de (*1R,2R,4S,5S,7s,9s*)-9-étil-7-[(*2S*)-2-fenil-3-hidroxipropanoil]oxi}-9-metil-3-oxa-9-azoniatriciclo[3.3.1.0^{2,4}]nonano

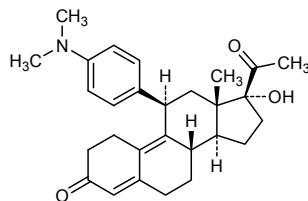


Recommended International Non Proprietary Names (Rec. INN): List 58
Dénominations communes internationales recommandées (DCI Rec.): Liste 58
Denominaciones Comunes Internacionales Recomendadas (DCI Rec.): Lista 58
(WHO Drug Information, Vol. 21, No. 3, 2007)

p. 260 **ulipristalum**

& 261 ulipristal
ulipristal
ulipristal

*replace the structure by the following
remplacer la structure par la suivante
sustitúyase la estructura por la siguiente*



Recommended International Non Proprietary Names (Rec. INN): List 60
Dénominations communes internationales recommandées (DCI Rec.): Liste 60
Denominaciones Comunes Internacionales Recomendadas (DCI Rec.): Lista 60
(WHO Drug Information, Vol. 22, No. 3, 2008)

p. 229 **conestatum alfa #**

conestat alfa
conestat alfa
conestat alfa

*replace the structure by the following
remplacer la structure par la suivante
sustitúyase la estructura por la siguiente*

```

NPNATSSSQ DPESLQDRGE GKVATTISKMLFVEPILEV SSLPTTNSTT 50
NSATKITANT TDEFTOPTT EPTTQPTIQP TQFTTQLPTD SPTQPTTGSF 100
CPGPVTLCSD LESHSTEAVL GDAIVDFSLK LYHAFSAMKK VETNMAFSFP 150
STASLLTQVL LGAGENIKTN LESLILSYPKD FTCVHQALKG FTTKGVTSVS 200
QIFHSPDLAI RDTFVNASRT LYSSSPRVLN NNSDANLELI NTWVAKNTNN 250
KISRLLDSLW SDTRIVLLNA IYLSAKWTT FDPKKTRMEP FHFKNNSVIKV 300
PMMSKKYKPV AHFIDQTLKA KVGGQLQLSHN LSLVILVPQN LKHRLLEDMEQ 350
ALSPSVFKAI MEKLEMMSKFQ PTLLTLPRIK VTTSQDMLSI MEKLEFFDFS 400
YDLNLCLGLTE DPDLQVSAMQ HQTVLELTET GVEAAAASAI SVARTLLLVE 450
VQQPFLFVWLW DQOHKFVVFM GRVYDPRW 478

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Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
101-406 108-183

Glycosylation sites / Sites de glycosylation / Posiciones de glicosilación
Asn-3 Thr-26 Ser-42 Asn-47 Thr-49 Asn-59 Thr-61
Thr-66 Thr-70 Thr-74 Asn-216 Asn-231 Asn-330

Recommended International Non Proprietary Names (Rec. INN): List 61
Dénominations communes internationales recommandées (DCI Rec.): Liste 61
Denominaciones Comunes Internacionales Recomendadas (DCI Rec.): Lista 61
(WHO Drug Information, Vol. 23, No. 1, 2009)

p. 67 **macitentanum**

macitentan

replace the chemical name by the following

N-[5-(4-bromophenyl)-6-{2-[(5-bromopyrimidin-2-yl)oxy]ethoxy}pyrimidin-4-yl]-N'-propylsulfuric diamide

Recommended International Non Proprietary Names (Rec. INN): List 62
Denominations communes internationales recommandées (DCI Rec.): Liste 62
Denominaciones Comunes Internacionales Recomendadas (DCI Rec.): Lista 62
(WHO Drug Information, Vol. 23, No. 3, 2009)

p. 263 & solanezumab #
 264

solanezumab *replace the description and the structure by the following ones*
 solanezumab *remplacer la description et la structure par les suivantes*
 solanezumab *sustitúyase la descripción y la estructura por las siguientes*

immunoglobulin G1-kappa, anti-[*Homo sapiens* amyloid-beta (Abeta) peptide soluble monomer], humanized monoclonal antibody;
 gamma1 heavy chain [humanized VH (*Homo sapiens* IGHV3-23*04 (87.60%) - (IGHD)-IGHJ4*01) [8.8.5] (1-112) -*Homo sapiens* IGHG1*01 (113-442)], (215-219')-disulfide with kappa light chain (1'-219') [humanized V-KAPPA (*Homo sapiens* IGKV2-30*01 (90.00%) -IGKJ1*01) [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113-219')]; (221-221":224-224")-bisdisulfide dimer

immunoglobuline G1-kappa, anti-[*Homo sapiens* amyloïde-β (Abeta) peptide monomère soluble], anticorps monoclonal humanisé;
 chaîne lourde gamma1 [VH humanisé (*Homo sapiens* IGHV3-23*04 (87.60%) - (IGHD)-IGHJ4*01) [8.8.5] (1-112) -*Homo sapiens* IGHG1*01 (113-442)], (215-219')-disulfure avec la chaîne légère kappa (1'-219') [V-KAPPA humanisé (*Homo sapiens* IGKV2-30*01 (90.00%) -IGKJ1*01) [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113-219')]; dimère (221-221":224-224")-bisdisulfure

inmunoglobulina G1-kappa, anti-[péptido amiloide-beta (Abeta) monomérico soluble de *Homo sapiens*], anticuerpo monoclonal humanizado;
 cadena pesada gamma1 [VH humanizado (*Homo sapiens* IGHV3-23*04 (87.60%) - (IGHD)-IGHJ4*01) [8.8.5] (1-112) -*Homo sapiens* IGHG1*01 (113-442)], (215-219')-disulfuro con la cadena ligera kappa (1'-219') [V-KAPPA humanizado (*Homo sapiens* IGKV2-30*01 (90.00%) -IGKJ1*01) [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113-219')]; dímero (221-221":224-224")-bisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada
 EVQLVESGGG LVQPGGSLRL SCAASGFTS RYSMSWVRQA PGKGLELVAQ 50
 INSVGNSTYY PDTVKGRFTI SRDNAKNTLY LQMSNSLAED TAVYVCASGD 100
 YWGQGTLVTV SSASTKGPSV FFLAPSSKST SGCTAALGCL VKDYFPEPV 150
 VSWNSGALTGS GVHTFPAVLQ SSGLYSLSSV VTVPSSSLGT QTYICVNHHK 200
 PSNTKVDKVV EPKSCDKTHT CPCPCAPELL GGPESVFLFPP KPKDTLMISR 250
 TPEVTCVVVD VSHEDPEVKF NWYVGVEVIR NARTKPRREEQ YNSTYRVVSV 300
 LTVLHQDWLN GKEYCKKVSN KALPAPIEKT ISAKKGQPRE PQVTLPSSR 350
 DELTKNQVSL TCLVKGFYPS DIAVEWESNG QPENNYKTTE PVLDSDGSFF 400
 LYSKLTVDKS RWQQGNVFSC SVMHEALHNH YTQKSLSLSP GK 442

Light chain / Chaîne légère / Cadena ligera
 DVVMTQSPLS LPVTLGQPAS ISCRSSQSLI YSDGNAYLHW FLQKPGQSPR 50
 LLIYKVSNRF SGVPDRFSGS GSGTDFTLKI SRVEAEDVGV YYCQSSTHVP 100
 WTFQGQTKEV IKRTVAAPSV FIFPPSDEQL KSCTASVCL LNNFYPREAK 150
 VQWKVDNALQ SGNSQESVTE QDSKDSTYSL SSTLTLSKAD YEKKHVYACE 200
 VTHQGLSSPV TKSFNRGEC 219

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-I 22"-96" 139"-195" 256"-316" 362"-420"
 22"-96" 139"-195" 256"-316" 362"-420"
 Intra-L 23"-93" 139"-199"
 23"-93" 139"-199"
 Inter-H-L 215-219' 215"-219"
 Inter-H-H 221-221" 224-224"

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
 H VH CDR2-IMGT N63:
 56, 56"
 H CH2 N84.4:
 292, 292"

Recommended International Non Proprietary Names (Rec. INN): List 64
Dénominations communes internationales recommandées (DCI Rec.): Liste 64
Denominaciones Comunes Internacionales Recomendadas (DCI Rec.): Lista 64
(WHO Drug Information, Vol. 23, No. 4, 2009)

p. 264 & **dalotuzumabum #**
 265

dalotuzumab *replace the description and the structure by the following ones*
 dalotuzumab *remplacer la description et la structure par les suivantes*
 dalotuzumab *sustitúyase la descripción y la estructura por las siguientes*

immunoglobulin G1-kappa, anti-[*Homo sapiens*] IGF1R (insulin-like growth factor 1 receptor, IGF1-R, IGF-1R, CD221), humanized monoclonal antibody; gamma1 heavy chain (1-447) [humanized VH (*Homo sapiens*)IGHV4-61*08 (86.90%) -(IGHD)-IGHJ4*01] [9.7.10] (1-117) -*Homo sapiens* IGHG1*03 (118-447)], (220-219')-disulfide with a kappa light chain (1'-219') [humanized V-KAPPA (*Homo sapiens*)IGKV2-29*03 (84.00%) -IGKJ1*01] [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113'-219')]; (226-226":229-229")-bisdisulfide dimer

immunoglobuline G1-kappa, anti-[*Homo sapiens*] IGF1R (récepteur du facteur de croissance 1 analogue à l'insuline (IGF1-R, IGF-1R, CD221)], anticorps monoclonal humanisé; chaîne lourde gamma1 (1-447) [VH humanisé (*Homo sapiens*)IGHV4-61*08 (86.90%) -(IGHD)-IGHJ4*01] [9.7.10] (1-117) -*Homo sapiens* IGHG1*03 (118-447)], (220-219')-disulfure avec la chaîne légère kappa (1'-219') [V-KAPPA humanisé (*Homo sapiens*)IGKV2-29*03 (84.00%) -IGKJ1*01] [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113'-219')]; dimère (226-226":229-229")-bisdisulfure

inmunoglobulina G1-kappa, anti-[*Homo sapiens*] IGF1R (receptor del factor de crecimiento similar a la insulina 1 (IGF1-R, IGF-1R, CD221)), anticuerpo monoclonal humanizado; cadena pesada gamma1 (1-447) [VH humanizado (*Homo sapiens*)IGHV4-61*08 (86.90%) -(IGHD)-IGHJ4*01] [9.7.10] (1-117) -*Homo sapiens* IGHG1*03 (118-447)], (220-219')-disulfuro con la cadena ligera kappa (1'-219') [V-KAPPA humanizado (*Homo sapiens*)IGKV2-29*03 (84.00%) -IGKJ1*01] [11.3.9] (1'-112') -*Homo sapiens* IGKC*01 (113'-219')]; dímero (226-226":229-229")-bisdisulfuro

Heavy chain / Chaîne lourde / Cadena pesada
 QVQLQESGPQ LVKPSETLSSL TCTVSGYSIT GGYLWNWIRQ PPGKGLEWIG 50
 YISYDGTNYY KPSLKDRVTI SRDTSKNQFS LKLSVVTAAAD TAVYYCARYG 100
 RVFFDYWGQQ TLTVVSSAST KGGSVPEFLAP SSKSTSGGTA ALGLCLVKDVF 150
 PEPVTVWSNS GALTSGVHTF PAVLQSSGLY SLSSVVTVP SSLGTQTYIC 200
 NVNHPKSNTK VDKRVEPKSC DKTHTCPFCP APELLGGPSV FLFPFPKPKDT 250
 LMISRTPETV CVVVVDVSHED PEVKFVNWYVD GVEVHNAAKTK FREEQYNSTY 300
 RRVSVLTVLH QDWLNKEYD CKVSNKALPA PIEKTISKAK GQPREPQVYT 350
 LPPSREEMTK NQVSLTCLVE GFYPSDIAVE WESNGOPENN YKTTTPVLDs 400
 DGSFFLYSKL TVDKSRWQG NVFSCSVMHE ALHNHYTQKS LSLSPGK 447

Light chain / Chaîne légère / Cadena ligera
 DIVMTQSPLS LPVTPGEPAAS ISCRSSQSIV HSNGNNTYLQW YLQKPGQSPQ 50
 LLIYKVSNRL YGVPDFRSFGS GSCTDFTELKI SRVEAEVGV YYCFQGSHVP 100
 WTFGGTKEVIE IKRTVAAPS FIFPPSDEQL KSGTASVCL LNNFYPREAK 150
 VQWKVDNALQ SGNSQESVTE QDSKDSTYSL SSTLTLSKAD YEKHKVYACE 200
 VTHQGLSSPV TKSFNRGEC 219

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 Intra-H 22-96 144-200 261-321 367-425
 22"-96" 144"-200" 261"-321" 367"-425"
 Intra-L 23"-93" 139"-199"
 23"-93" 139"-199"
 Inter-H-L 220-219" 220"-219"
 Inter-H-H 226-226" 229-229"

N-glycosylation sites / Sites de N-glycosylation / Posiciones de N-glicosilación
 H CH2 N84.4
 297, 297"

Recommended International Non Proprietary Names (Rec. INN): List 67
Dénominations communes internationales recommandées (DCI Rec.): Liste 67
Denominaciones Comunes Internacionales Recomendadas (DCI Rec.): Lista 67
(WHO Drug Information, Vol. 26, No. 1, 2012)

p. 70 **lipogfilgrastim #**

lipogfilgrastim *replace the description and the structure by the following ones*
 lipogfilgrastim *remplacer la description et la structure par les suivantes*
 lipogfilgrastim *sustitúyase la descripción y la estructura por las siguientes*

pegylated granulocyte colony stimulating factor (human short isoform);
 $O^{3,137}-\{[3,5-dideoxy-5-[(N-[\omega-methoxypoly(oxyethylene)]carbonyl)glycyl]amino]-D-glycero-\alpha-D-galacto-non-2-ulopyranosylonic acid\}-(2\rightarrow6)-2-(acetylamino)-2-deoxy-\alpha-D-galactopyranosyl]-des-(37-39)-[1-methionine]human granulocyte colony-stimulating factor (G-CSF, pluripoietin)$

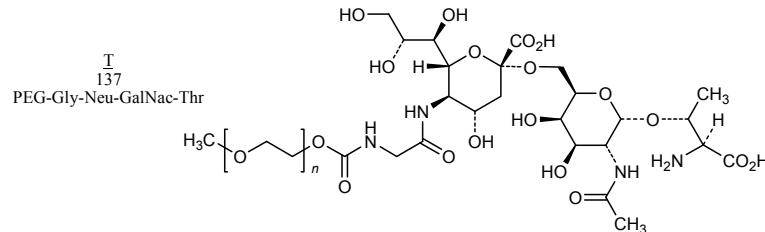
facteur de stimulation de colonie de granulocytes (isoforme court humain) pégylé;
 $O^{3,137}-\{[acide\ 3,5-didésoxy-5-[(N-[\omega-méthoxypoli(oxyéthylène)]carbonyl)glycyl]amino]-D-glycéro-\alpha-D-galacto-non-2-ulopyranosylonique\}-(2\rightarrow6)-2-(acétylamino)-2-déoxy-\alpha-D-galactopyranosyl]-dès-(37-39)-[1-méthionine]facteur humain de stimulation de colonie de granulocytes (G-CSF, pluripoétine)$

factor de estimulación de colonias de granulocitos (isoformo corto humano) pegilado;
 $O^{3,137}-\{[ácido\ 3,5-didesoxi-5-[(N-[\omega-metoxipoli(oxietileno)]carbonil)glicil]amino]-D-glicero-\alpha-D-galacto-non-2-ulopiranosilónico\}-(2\rightarrow6)-2-(acetilamino)-2-desoxi-\alpha-D-galactopiranosil]-dés-(37-39)-[1-méthionina]factor humano de estimulación de colonias de granulocitos (G-CSF, pluripoyetina)$

MTPLGPASSL PQSFLLKCLE QVRKIQGDGA ALQEKL---C ATYKLCHPEE 50
 LVLIGHSLGI PWAPLSSCPS QALQLAGCLS QLHSGLFLYQ GLLQALEGIS 100
 PELGPTLDL QLDVADFTT IWQQMEELGM APALQFTQGA MPAFASAFQR 150
 RAGGVLVASH LQSFLVEVSYR VLRHLAQP 178

Disulfide bridges location / Position des ponts disulfure / Posiciones de los puentes disulfuro
 40-46 68-78

Modified residues / Résidu modifié / Residuo modificado



Electronic structure available on Mednet: <http://mednet.who.int/>
Structure électronique disponible sur Mednet: <http://mednet.who.int/>
Estructura electrónica disponible en Mednet: <http://mednet.who.int/>

Procedure and Guiding Principles / Procédure et Directives / Procedimientos y principios generales

The text of the *Procedures for the Selection of Recommended International Nonproprietary Names for Pharmaceutical Substances and General Principles for Guidance in Devising International Nonproprietary Names for Pharmaceutical Substances* will be reproduced in proposed INN lists only.

Les textes de la *Procédure à suivre en vue du choix de dénominations communes internationales recommandées pour les substances pharmaceutiques* et des *Directives générales pour la formation de dénominations communes internationales applicables aux substances pharmaceutiques* seront publiés seulement dans les listes des DCI proposées.

El texto de los *Procedimientos de selección de denominaciones comunes internacionales recomendadas para las sustancias farmacéuticas* y de los *Principios generales de orientación para formar denominaciones comunes internacionales para sustancias farmacéuticas* aparece solamente en las listas de DCI propuestas.