

# STATEMENT ON A NONPROPRIETARY NAME ADOPTED BY THE USAN COUNCIL

USAN (fg-27)

CAMIDANLUMAB TESIRINE

PRONUNCIATION

kam" i dan' lue mab tes ir' een

THERAPEUTIC CLAIM

Treatment of hematologic cancers

CHEMICAL NAMES

1. Immunoglobulin G1, anti-(human interleukin 2 receptor  $\alpha$  chain)(human HuMax-TAC heavy chain), human HuMax-TAC light chain, dimer, thioether with *N*-[31-(2,5-dihydro-2,5-dioxo-1*H*-pyrrol-1-yl)-1,29-dioxo-4,7,10,13,16,19,22,25-octaoxa-28-azahentriacont-1-yl]-*L*-valyl-*N*-[4-[[[(11*S*,11*aS*)-8-[[5-[[[(11*aS*)-5,11*a*-dihydro-7-methoxy-2-methyl-5-oxo-1*H*-pyrrolo[2,1-*c*][1,4]benzodiazepin-8-yl]oxy]pentyl]oxy]-11,11*a*-dihydro-11-hydroxy-7-methoxy-2-methyl-5-oxo-1*H*-pyrrolo[2,1-*c*][1,4]benzodiazepine-10(5*H*)-yl]carbonyl]oxy]methyl]phenyl]-*L*-alaninamide
2. Immunoglobulin G1-kappa, anti-[*Homo sapiens* Interleukin-2 receptor subunit alpha (TAC antigen, p55, CD25 antigen)], *Homo sapiens* monoclonal antibody conjugated to an average of two molecules of tesirine;  $\gamma$ 1 heavy chain (1-445) [*Homo sapiens* VH (IGHV1-69\*02 (95%) –(IGHD)-IGHJ4\*01 (93%)) [8.8] (1-115) -*Homo sapiens* IGHG1\*03 (116-445)], (218-214')-disulfide with  $\kappa$  light chain (1'-214') [*Homo sapiens* V-KAPPA (IGKV3-20\*01 (99%) –IGKJ4\*01)[6.3.9] (1'-107') -*Homo sapiens* IGKC\*01 (108'-214')]; dimer (224-224":227-227")-bisdisulfide, an average of two cysteines are *S*-linked to (1<sup>11*a*</sup>*S*,9<sup>11</sup>*S*,9<sup>11*a*</sup>*S*,16*S*,19*S*,52<sup>3</sup>*RS*)-9<sup>11</sup>-hydroxy-1<sup>7</sup>, 9<sup>7</sup>-dimethoxy-1<sup>2</sup>,9<sup>2</sup>,16-trimethyl-1<sup>5</sup>,9<sup>5</sup>,10,15,18,21,49,52<sup>2</sup>-52<sup>5</sup>-nonaoxo-19-(propan-2-yl)-1<sup>5</sup>,1<sup>11*a*</sup>,9<sup>11</sup>,9<sup>11*a*</sup>-tetrahydro-1<sup>1</sup>*H*,9<sup>1</sup>*H*,9<sup>5</sup>*H*-2,8,11,24,27,30,33,36,39,42,45-undecaoxa-14,17,20,48-tetraaza-1(8),9(8,10)-bis(pyrrolo[2,1-*c*][1,4]benzodiazepine)-52(1)-pyrrolidina-13(1,4)benzenadopentacontaphan-52<sup>3</sup>-yl

## STRUCTURAL FORMULA

Heavy chain

QVQLVQSGAE	VKKPGSSVKV	SCKASGGTFS	RYIINWVRQA	PGQGLEWMGR	50
IIPILGVENY	AQKFQGRVTI	TADKSTSTAY	MELSSLRSED	TAVYYCARKD	100
WFDYWGQGTL	VTVSSASTKG	PSVFLAPSS	KSTSGGTAAL	GCLVKDYFPE	150
PVTVSWNSGA	LTSGVHTFPA	VLQSSGLYSL	SSVVTVPSSS	LGTQTYICNV	200
NHKPSNTKVD	KRVEPKSCDK	THTCPPCPAP	ELLGGPSVFL	FPPKPKDTLM	250
ISRTPEVTCV	VVDVSHEDPE	VKFNWYVDGV	EVHNAKTKPR	EEQYNSTYRV	300
VSVLTVLHQD	WLNQKEYKCK	VSNKALPAPI	EKTIISKAKGQ	PREPQVYTLP	350
PSREEMTKNQ	VSLTCLVKGF	YPSDIAVEWE	SNGQPENNYK	TTPPVLDSDG	400
SFFLYSKLTV	DKSRWQQGNV	FSCSVMHEAL	HNHYTQKSLS	LSPGK	445

Light chain

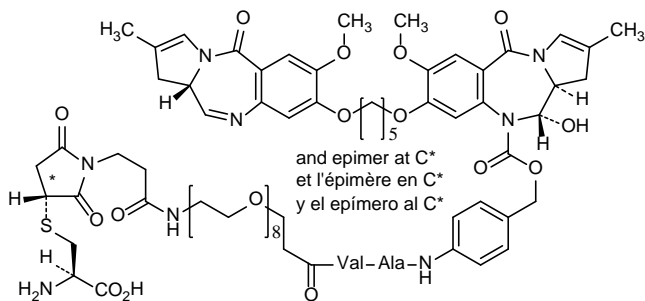
EIVLTQSPGT	LSLSPGERAT	LSCRASQSVS	SYLAWYQQKP	GQAPRLLIYG	50'
ASSRATGIPD	RFSGSGSGTD	F <del>L</del> T <del>L</del> ISRLEP	EDFAVYYCQQ	YGSSPLTFGG	100'
GTKVEIKRTV	AAPSVFIFPP	SDEQLKSGTA	SVVCLLN <del>N</del> FY	PREAKVQWKV	150'
DNALQSGNSQ	ESVTEQDSK	STYLSSTLT	LSKADYEKHK	VYACEVTHQG	200'
LSSPVTKSFN	RGEC				214'

Disulfide bridges

22-96	22"-96"	23'-88'	23'''-88'''	134'-194'	134'''-194'''	142-198	142"-198"
214'-218*	214'''-218'''*	224-224**	227-227**	259-319	259"-319"	365-423	365'''-423'''

\* one or two of the inter-chain disulfide bridges are not present, an average of two Cysteines are S-substituted with tesirine C.

Modified residues (C)



Glycosylation sites (N)

Asn-295	Asn-295"
---------	----------

MOLECULAR FORMULA

C<sub>6548</sub>H<sub>10150</sub>N<sub>1732</sub>O<sub>2038</sub>S<sub>42</sub>

MOLECULAR WEIGHT

147.1 kDa

TRADEMARK

None as yet

SPONSOR

ADC Therapeutics SA

CODE DESIGNATIONS

ADCT-301

CAS REGISTRY NUMBER

1853239-04-9

UNII

LYJ1AEJ9YH

WHO NUMBER

10592

gbk