

LISTADO DE SECUENCIAS

<110> Arvinas Operations, Inc.

<120> COMPUESTOS Y MÉTODOS PARA LA DEGRADACIÓN DIRIGIDA DEL RECEPTOR DE ANDRÓGENOS

<130> ARVN-007/001WO 331216-2140

<150> 63/089,497

<151> 2020-10-08

<150> 63/032,473

<151> 2020-05-29

<150> 62/950,815

<151> 2019-12-19

<160> 1

<170> PatentIn versión 3.5

<210> 1

<211> 920

<212> PRT

<213> Homo sapiens

<400> 1

Met	Glu	Val	Gln	Leu	Gly	Leu	Gly	Arg	Val	Tyr	Pro	Arg	Pro	Pro	Ser
1				5					10					15	

Lys	Thr	Tyr	Arg	Gly	Ala	Phe	Gln	Asn	Leu	Phe	Gln	Ser	Val	Arg	Glu
			20					25					30		

Val	Ile	Gln	Asn	Pro	Gly	Pro	Arg	His	Pro	Glu	Ala	Ala	Ser	Ala	Ala
		35					40					45			

Pro	Pro	Gly	Ala	Ser	Leu	Leu	Leu	Leu	Gln	Gln	Gln	Gln	Gln	Gln	Gln
	50					55				60					

Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln
65					70				75					80	

Glu	Thr	Ser	Pro	Arg	Gln	Gln	Gln	Gln	Gln	Gln	Gly	Glu	Asp	Gly	Ser
				85					90					95	

Pro	Gln	Ala	His	Arg	Arg	Gly	Pro	Thr	Gly	Tyr	Leu	Val	Leu	Asp	Glu
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110

Asp Thr Ala Glu Tyr Ser Pro Phe Lys Gly Gly Tyr Thr Lys Gly Leu
305 310 315 320

Glu Gly Glu Ser Leu Gly Cys Ser Gly Ser Ala Ala Ala Gly Ser Ser
325 330 335

Gly Thr Leu Glu Leu Pro Ser Thr Leu Ser Leu Tyr Lys Ser Gly Ala
340 345 350

Leu Asp Glu Ala Ala Ala Tyr Gln Ser Arg Asp Tyr Tyr Asn Phe Pro
355 360 365

Leu Ala Leu Ala Gly Pro Pro Pro Pro Pro Pro Pro Pro His Pro His
370 375 380

Ala Arg Ile Lys Leu Glu Asn Pro Leu Asp Tyr Gly Ser Ala Trp Ala
385 390 395 400

Ala Ala Ala Ala Gln Cys Arg Tyr Gly Asp Leu Ala Ser Leu His Gly
405 410 415

Ala Gly Ala Ala Gly Pro Gly Ser Gly Ser Pro Ser Ala Ala Ala Ser
420 425 430

Ser Ser Trp His Thr Leu Phe Thr Ala Glu Glu Gly Gln Leu Tyr Gly
435 440 445

Pro Cys Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly
450 455 460

Gly Gly Gly Gly Gly Gly Gly Gly Gly Glu Ala Gly Ala Val Ala Pro
465 470 475 480

Tyr Gly Tyr Thr Arg Pro Pro Gln Gly Leu Ala Gly Gln Glu Ser Asp
485 490 495

Phe Thr Ala Pro Asp Val Trp Tyr Pro Gly Gly Met Val Ser Arg Val
500 505 510

Pro Tyr Pro Ser Pro Thr Cys Val Lys Ser Glu Met Gly Pro Trp Met
515 520 525

Asp Ser Tyr Ser Gly Pro Tyr Gly Asp Met Arg Leu Glu Thr Ala Arg

530

535

540

Asp His Val Leu Pro Ile Asp Tyr Tyr Phe Pro Pro Gln Lys Thr Cys
 545 550 555 560

Leu Ile Cys Gly Asp Glu Ala Ser Gly Cys His Tyr Gly Ala Leu Thr
 565 570 575

Cys Gly Ser Cys Lys Val Phe Phe Lys Arg Ala Ala Glu Gly Lys Gln
 580 585 590

Lys Tyr Leu Cys Ala Ser Arg Asn Asp Cys Thr Ile Asp Lys Phe Arg
 595 600 605

Arg Lys Asn Cys Pro Ser Cys Arg Leu Arg Lys Cys Tyr Glu Ala Gly
 610 615 620

Met Thr Leu Gly Ala Arg Lys Leu Lys Lys Leu Gly Asn Leu Lys Leu
 625 630 635 640

Gln Glu Glu Gly Glu Ala Ser Ser Thr Thr Ser Pro Thr Glu Glu Thr
 645 650 655

Thr Gln Lys Leu Thr Val Ser His Ile Glu Gly Tyr Glu Cys Gln Pro
 660 665 670

Ile Phe Leu Asn Val Leu Glu Ala Ile Glu Pro Gly Val Val Cys Ala
 675 680 685

Gly His Asp Asn Asn Gln Pro Asp Ser Phe Ala Ala Leu Leu Ser Ser
 690 695 700

Leu Asn Glu Leu Gly Glu Arg Gln Leu Val His Val Val Lys Trp Ala
 705 710 715 720

Lys Ala Leu Pro Gly Phe Arg Asn Leu His Val Asp Asp Gln Met Ala
 725 730 735

Val Ile Gln Tyr Ser Trp Met Gly Leu Met Val Phe Ala Met Gly Trp
 740 745 750

Arg Ser Phe Thr Asn Val Asn Ser Arg Met Leu Tyr Phe Ala Pro Asp
755 760 765

Leu Val Phe Asn Glu Tyr Arg Met His Lys Ser Arg Met Tyr Ser Gln
770 775 780

Cys Val Arg Met Arg His Leu Ser Gln Glu Phe Gly Trp Leu Gln Ile
785 790 795 800

Thr Pro Gln Glu Phe Leu Cys Met Lys Ala Leu Leu Leu Phe Ser Ile
805 810 815

Ile Pro Val Asp Gly Leu Lys Asn Gln Lys Phe Phe Asp Glu Leu Arg
820 825 830

Met Asn Tyr Ile Lys Glu Leu Asp Arg Ile Ile Ala Cys Lys Arg Lys
835 840 845

Asn Pro Thr Ser Cys Ser Arg Arg Phe Tyr Gln Leu Thr Lys Leu Leu
850 855 860

Asp Ser Val Gln Pro Ile Ala Arg Glu Leu His Gln Phe Thr Phe Asp
865 870 875 880

Leu Leu Ile Lys Ser His Met Val Ser Val Asp Phe Pro Glu Met Met
885 890 895

Ala Glu Ile Ile Ser Val Gln Val Pro Lys Ile Leu Ser Gly Lys Val
900 905 910

Lys Pro Ile Tyr Phe His Thr Gln
915 920