

# SEQUENCE LISTING

<110> Faron Pharmaceuticals Oy  
 <120> Use of an agent capable of binding to CLEVER-1 in immune activation and methods based thereon  
 <130> AP106793  
 <150> FI20165336  
 <151> 2016-04-18  
 <160> 7  
 <170> PatentIn version 3.5  
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 <212> PRT  
 <213> Homo sapiens  
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Gln Glu Ile Thr Val Thr Phe Asn Gln Phe Thr Lys  
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<400> 7

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35 40 45

Gly Gln Val Leu Phe Lys Gly Cys Asp Val Lys Thr Thr Phe Val Thr  
50 55 60

His Val Pro Cys Thr Ser Cys Ala Ala Ile Lys Lys Gln Thr Cys Pro  
65 70 75 80

Ser Gly Trp Leu Arg Glu Leu Pro Asp Gln Ile Thr Gln Asp Cys Arg  
85 90 95

Tyr Glu Val Gln Leu Gly Gly Ser Met Val Ser Met Ser Gly Cys Arg  
100 105 110

Arg Lys Cys Arg Lys Gln Val Val Gln Lys Ala Cys Cys Pro Gly Tyr  
115 120 125

Trp Gly Ser Arg Cys His Glu Cys Pro Gly Gly Ala Glu Thr Pro Cys  
130 135 140

Asn Gly His Gly Thr Cys Leu Asp Gly Met Asp Arg Asn Gly Thr Cys  
145 150 155 160

Val Cys Gln Glu Asn Phe Arg Gly Ser Ala Cys Gln Glu Cys Gln Asp  
165 170 175

Pro Asn Arg Phe Gly Pro Asp Cys Gln Ser Val Cys Ser Cys Val His  
180 185 190

Gly Val Cys Asn His Gly Pro Arg Gly Asp Gly Ser Cys Leu Cys Phe  
Sivu 2

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Ala	Gly	Tyr	Thr	Gly	Pro	His	Cys	Asp	Gln	Glu	Leu	Pro	Val	Cys	Gln
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Glu	Leu	Arg	Cys	Pro	Gln	Asn	Thr	Gln	Cys	Ser	Ala	Glu	Ala	Pro	Ser
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Cys	Arg	Cys	Leu	Pro	Gly	Tyr	Thr	Gln	Gln	Gly	Ser	Glu	Cys	Arg	Ala
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Pro	Asn	Pro	Cys	Trp	Pro	Ser	Pro	Cys	Ser	Leu	Leu	Ala	Gln	Cys	Ser
			260					265					270		
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Gly	Asp	Gly	Met	Val	Cys	Leu	Pro	Lys	Asp	Pro	Cys	Thr	Asp	Asn	Leu
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Gln	Ala	Phe	Cys	Thr	Cys	Arg	Pro	Gly	Leu	Val	Ser	Ile	Asn	Ser	Asn
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Ala	Ser	Ala	Gly	Cys	Phe	Ala	Phe	Cys	Ser	Pro	Phe	Ser	Cys	Asp	Arg
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Ser	Ala	Thr	Cys	Gln	Val	Thr	Ala	Asp	Gly	Lys	Thr	Ser	Cys	Val	Cys
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Arg	Glu	Ser	Glu	Val	Gly	Asp	Gly	Arg	Ala	Cys	Tyr	Gly	His	Leu	Leu
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His	Glu	Val	Gln	Lys	Ala	Thr	Gln	Thr	Gly	Arg	Val	Phe	Leu	Gln	Leu
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Arg	Val	Ala	Val	Ala	Met	Met	Asp	Gln	Gly	Cys	Arg	Glu	Ile	Leu	Thr
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Thr	Ala	Gly	Pro	Phe	Thr	Val	Leu	Val	Pro	Ser	Val	Ser	Ser	Phe	Ser
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Ser	Arg	Thr	Met	Asn	Ala	Ser	Leu	Ala	Gln	Gln	Leu	Cys	Arg	Gln	His
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Ile	Ile	Ala	Gly	Gln	His	Ile	Leu	Glu	Asp	Thr	Arg	Thr	Gln	Gln	Thr
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Arg	Arg	Trp	Trp	Thr	Leu	Ala	Gly	Gln	Glu	Ile	Thr	Val	Thr	Phe	Asn
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Gln Phe Thr Lys Tyr Ser Tyr Lys Tyr Lys Asp Gln Pro Gln Gln Thr  
 485 490 495

Phe Asn Ile Tyr Lys Ala Asn Asn Ile Ala Ala Asn Gly Val Phe His  
 500 505 510

Val Val Thr Gly Leu Arg Trp Gln Ala Pro Ser Gly Thr Pro Gly Asp  
 515 520 525

Pro Lys Arg Thr Ile Gly Gln Ile Leu Ala Ser Thr Glu Ala Phe Ser  
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Arg Phe Glu Thr Ile Leu Glu Asn Cys Gly Leu Pro Ser Ile Leu Asp  
 545 550 555 560

Gly Pro Gly Pro Phe Thr Val Phe Ala Pro Ser Asn Glu Ala Val Asp  
 565 570 575

Ser Leu Arg Asp Gly Arg Leu Ile Tyr Leu Phe Thr Ala Gly Leu Ser  
 580 585 590

Lys Leu Gln Glu Leu Val Arg Tyr His Ile Tyr Asn His Gly Gln Leu  
 595 600 605

Thr Val Glu Lys Leu Ile Ser Lys Gly Arg Ile Leu Thr Met Ala Asn  
 610 615 620

Gln Val Leu Ala Val Asn Ile Ser Glu Glu Gly Arg Ile Leu Leu Gly  
 625 630 635 640

Pro Glu Gly Val Pro Leu Gln Arg Val Asp Val Met Ala Ala Asn Gly  
 645 650 655

Val Ile His Met Leu Asp Gly Ile Leu Leu Pro Pro Thr Ile Leu Pro  
 660 665 670

Ile Leu Pro Lys His Cys Ser Glu Glu Gln His Lys Ile Val Ala Gly  
 675 680 685

Ser Cys Val Asp Cys Gln Ala Leu Asn Thr Ser Thr Cys Pro Pro Asn  
 690 695 700

Ser Val Lys Leu Asp Ile Phe Pro Lys Glu Cys Val Tyr Ile His Asp  
 705 710 715 720

Pro Thr Gly Leu Asn Val Leu Lys Lys Gly Cys Ala Ser Tyr Cys Asn  
 725 730 735

Gln Thr Ile Met Glu Gln Gly Cys Cys Lys Gly Phe Phe Gly Pro Asp  
 740 745 750

Cys Thr Gln Cys Pro Gly Gly Phe Ser Asn Pro Cys Tyr Gly Lys Gly  
 755 760 765  
 Asn Cys Ser Asp Gly Ile Gln Gly Asn Gly Ala Cys Leu Cys Phe Pro  
 770 775 780  
 Asp Tyr Lys Gly Ile Ala Cys His Ile Cys Ser Asn Pro Asn Lys His  
 785 790 795 800  
 Gly Glu Gln Cys Gln Glu Asp Cys Gly Cys Val His Gly Leu Cys Asp  
 805 810 815  
 Asn Arg Pro Gly Ser Gly Gly Val Cys Gln Gln Gly Thr Cys Ala Pro  
 820 825 830  
 Gly Phe Ser Gly Arg Phe Cys Asn Glu Ser Met Gly Asp Cys Gly Pro  
 835 840 845  
 Thr Gly Leu Ala Gln His Cys His Leu His Ala Arg Cys Val Ser Gln  
 850 855 860  
 Glu Gly Val Ala Arg Cys Arg Cys Leu Asp Gly Phe Glu Gly Asp Gly  
 865 870 875 880  
 Phe Ser Cys Thr Pro Ser Asn Pro Cys Ser His Pro Asp Arg Gly Gly  
 885 890 895  
 Cys Ser Glu Asn Ala Glu Cys Val Pro Gly Ser Leu Gly Thr His His  
 900 905 910  
 Cys Thr Cys His Lys Gly Trp Ser Gly Asp Gly Arg Val Cys Val Ala  
 915 920 925  
 Ile Asp Glu Cys Glu Leu Asp Met Arg Gly Gly Cys His Thr Asp Ala  
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 Leu Cys Ser Tyr Val Gly Pro Gly Gln Ser Arg Cys Thr Cys Lys Leu  
 945 950 955 960  
 Gly Phe Ala Gly Asp Gly Tyr Gln Cys Ser Pro Ile Asp Pro Cys Arg  
 965 970 975  
 Ala Gly Asn Gly Gly Cys His Gly Leu Ala Thr Cys Arg Ala Val Gly  
 980 985 990  
 Gly Gly Gln Arg Val Cys Thr Cys Pro Pro Gly Phe Gly Gly Asp Gly  
 995 1000 1005  
 Phe Ser Cys Tyr Gly Asp Ile Phe Arg Glu Leu Glu Ala Asn Ala  
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His	Phe	Ser	Ile	Phe	Tyr	Gln	Trp	Leu	Lys	Ser	Ala	Gly	Ile	Thr
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Leu	Pro	Ala	Asp	Arg	Arg	Val	Thr	Ala	Leu	Val	Pro	Ser		
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