

レポートの冒頭に、検索の詳細
(検索タイプやパラメータなど)が
表示される

Query Input

Result Name: p53 related gene
Program: BLASTn
Subsets: Patents
Non-patents
EST
STS
GSS
HTGS
Other GenBank (R)

Return Sequence Length: 50 or fewer bases
51 - 500 bases
501 or more bases

Date Range: Complete Database

Low Complexity Filtering: Off
Max No. of Answers: 1,000
Expectation Value: 10
Word Size: 11
Open Gap Cost: 5
Extend Gap Cost: 2
Penalty for Mismatch: -3
Reward for Match: 1

← 検索パラメータ

Sequence:

gctcccagaa tgccagaggc tgctccccc gtggcccctg caccagcgcac tcctacaccg
gcgcccctg caccagcccc ctccctggccc ctgtcatctt ctgtcccctc ccagaaaacc
taccagggca gctacggttt ccgtctgggc ttcttgcatc ctgggacagc caagtctgtg
acttgcaagt actcccctgc cctcaacaag atgttttgcc aactggccaa gacctgccct
gtgcagctgt gggttgattc cacaccccc cccggcaccg ggtccgcgc catggccatc
tac

← 配列質問式

Sequence Identifier:

Result Summary

Result Name: p53 related gene
Program: BLASTn
Creation Date/Time: 18/08/21 11:51
Unique Sequences: 1,000
Total Sequences: 1,383

Result Set Sequences

← 回答全件の CAS RN®, 名称, スコア値, 期待値

Sequence	Score	E-Value
(1515955-91-5) 30: PN: US8613907 SEQID: 30 unclaimed DNA	601	2e-168
(457009-82-4) GenBank BD135657: Observation alley for exp...	601	2e-168
:		
(256805-43-3) GenBank AC023805: Mus musculus clone RP23-4...	105	3e-19
(140765-91-9) GenBank X00878: Murine gene fragment for ce...	105	3e-19

CAplus ファイルの例
 BIB HITRN で表示
 Use format : Standard
 (デフォルトで選択されたフィールドを利用)

L2 ANSWER 1 OF 3 CAPLUS COPYRIGHT 2018 ACS on STN

[PatentPak PDF](#)

TI Methods and compositions for preventing or treating cancer

AN 2017:594861 CAPLUS [Full-text](#)

DT Patent

LA English

IN Schiffman, Joshua; Schroeder, Avi; Abegglen, Lisa

PA University of Utah Research Foundation, USA; Technion Research & Development Foundation Limited

DN 166:435154

FAN.CNT 1

SO PCT Int. Appl., 49pp.

CODEN: PIXXD2

PPPI

PATENT NO.	KIND	DATE	LANGUAGE	PatentPak
WO 2017062726	A1	20170413	English	PDF
KR 2018086188	A	20180730	Korean	PDF

PI

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2017062726	A1	20170413	WO 2016-US55921	20161007
W:			AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BN, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CZ, DE, DJ, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IR, IS, JP, KE, KG, KN, KP, KR, KW, KZ, LA, LC, LK, LR, LS, LU, LY, MA, MD, ME, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PA, PE, PG, PH, PL, PT, QA, RO, RS, RU, RW, SA, SC, SD, SE, SG, SK, SL, SM, ST, SV, SY, TH, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, ZA, ZM, ZW	
RW:			AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, KM, ML, MR, NE, SN, TD, TG, BW, GH, GM, KE, LR, LS, MW, MZ, NA, RW, SD, SL, ST, SZ, TZ, UG, ZM, ZW, AM, AZ, BY, KG, KZ, RU, TJ, TM	
CA 3001054	A1	20170413	CA 2016-3001054	20161007
AU 2016335701	A1	20180510	AU 2016-335701	20161007
IL 258512	A	20180531	IL 2016-258512	20161007
KR 2018086188	A	20180730	KR 2018-7012840	20161007
EP 3359197	A1	20180815	EP 2016-854394	20161007
R:			AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LI, LT, LU, LV, MC, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR, BA, ME	
IN 201817016788	A	20180810	IN 2018-17016788	20180503
PRAI US 2015-62239103	P	20151008		
US 2016-62379179	P	20160824		
WO 2016-US55921	W	20161007		

IT 2094085-15-9

RL: BSU (Biological study, unclassified); PRP (Properties); THU
(Therapeutic use); BIOL (Biological study); USES (Uses)
(nucleotide sequence; methods and compns. for preventing or treating
cancer)

2094085-15-9

BLAST® Alignment Data

Length = 1129 Score = 135 Expect = 4e-28

Score = 135 Expect = 4e-28

Identities = 198/240 (82%) Gaps = 1/240 (0%)

Strand = Plus/Plus

Query: 64 GCCCCTGCACCAGCCCCCTCCTGGCCCTGTCATCTTCTGTCCCTTCCCAGAAAA 118

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Subject: 202 GCCCCTACACTAGCCACCTCCTGGATGCTGTCATCCTCTGTCCCTTCTCAGAAGA 256

Query: 119 CCTACCAGGGCAGCTACGGTTTCCGTCTGGGCTTCTTGCACTTCTGGGACAGCCAA 173

||| || | || ||| |||| | ||||| | ||||| | ||||| | ||||| |

Subject: 257 CTGCCCAG-CACCTATCGTTTCTGTCTGGGCTTCTTGCACTTCTGGGACAGCCAA 310

Query: 174 GTCTGTGACTTGCACGTACTCCCCTGCCCTCAACAAGATGTTTTGCCAACTGGCC 228

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Subject: 311 GTCCGTACCTACACATACTCCCCTGAACTTAACATGCTGTTTTGCCAGCTGGCA 365

Query: 229 AAGACCTGCCCTGTGCAGCTGTGGTTGATTCCACACCCCGCCCGGCACCCGCG 283

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Subject: 366 AAGGCCTGCCAGTGCAGCTGTGGTCACTCAACACCCCGCCAGCACCTGTG 420

Query: 284 TCCGCGCCATGGCCATCTAC 303

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Subject: 421 TTCACACCATGGCCATCTAC 440

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CAS RN® に対応するアライメントを
レポートに含めることができる