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## Primer-BLAST

» **JOB ID:iYNXuWgmZY5CsGC1bdVEhxfOVbU63U6oOw**

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Primer-BLAST Results

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**Input PCR template**

none

### Specificity of primers

Target templates were found in selected database: RefSeq Representative Genome Database (Organism limited to Homo sapiens)

### Other reports

[Search Summary](#)

Detailed primer reports

You can re-search for specific primers by accepting some of the unintended targets, check the box(es) next to the ones you accept and try again to re-search for specific primers

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## Primer pair 1

	Sequence (5'->3')	Length	Tm	GC%	Self complementarity	Self 3' complementarity
<b>Forward primer</b>	ATGGGAAGACCCAGTTAGCGT	21	61.18	52.38	4.00	0.00
<b>Reverse primer</b>	GAGGAAGGTGGCAAGAGGTGTAGA	24	63.66	54.17	3.00	2.00

Products on intended targets

Products on allowed targets

Products on allowed transcript variants

Products on potentially unintended templates

Products on target templates

>[NC\\_000012.12](#) Homo sapiens chromosome 12, GRCh38.p14 Primary Assembly

product length = 345

Features associated with this product:

[corrinoid adenosyltransferase precursor](#)

[corrinoid adenosyltransferase isoform x3](#)

Forward primer	1	ATGGGAAGACCCAGTTAGCGT	21
Template	109557209	.....	109557189
Reverse primer	1	GAGGAAGGTGGCAAGAGGTGTAGA	24
Template	109556865	.....	109556888

If you want to allow any of the unintended targets, check the box(es) next to the ones you accept and try again to re-search for specific primers

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