## **Certificate of Analysis**

# **.:: REACTION** .:::: BIOLOGY

## ProQinase<sup>™</sup> TESTPRODUKT LOT002

### XXX standard example full name

#### **Recombinant Human Active Protein Kinase**

HGNC Symbol: KIN1

Synonyms: Kanase, Konase

Product No.: 0000-0000-1

Lot: 000

**Description:** Human Kinase, C/N/internal fragment, amino acids M- $_1$ -X $_{1000}$  (as in NCBI/Protein entry NP\_000000.0), N-terminal GST-HIS<sub>6</sub> fusion protein with a Thrombin/3C cleavage site, expressed in Sf9 insect cells/E.coli

Product identity: Kinase Lot 000, was confirmed as Kinase by mass spectroscopy LC-ESI-MS/MS

Theoretical MW<sub>Fusion Protein</sub>: 00,000 Da

Expression host: Sf9 insect cells/E.coli

Purification: GST-Affinity Chromatography Immobilized Metal Affinity Chromatography

Activation: This kinase was not activated by special procedures / With KinaseY / in vitro auto activation

Storage buffer: 50 mM HEPES pH 7.5, 100 mM NaCl, 5 mM DTT, 15 mM reduced glutathione, 20 % glycerol

Storage temperature: -80°C

For complete recovery, mix well and spin before use. Product must not be stored in diluted solutions, aliquots below 10µl are not advisable. Avoid repeated freeze-thaw cycles!

**Protein concentration:** 0.000 μg/μl (Bradford method using BSA [Sigma, cat# A-7638, Lot 79H7641] as standard protein)

**Biochemical Parameters:** 

Specific kinase activity (P\_i transfer): 0 pmol/µg\*min ATP-K\_M: 0 µM

Additional assay technology:

Kinase Lot 000 was also successfully tested by Reaction Biology for the use with the ADP-Glo Kinase assay from Promega ADP-Glo assay conditions may vary from radiometric assay conditions, please inquire for assay details

Kinase Lot 000: Coomassie stain

Kinase Lot 000: Determination of  $V_{\text{max}}$  and  $K_{\text{M}}$  value for ATP

 Assay conditions:

 60 mM HEPES-NaOH, pH 7.5
 3 mM MgCl<sub>2</sub>
 3 mM MnCl<sub>2</sub>
 3 µM Na-orthovanadate
 1.2 mM DTT
 50 µg/ml PEG<sub>20.000</sub>
 ATP (variable)
 Substrate: Example 0 µg/ml Kinase: 0 µg/ml

Filter binding assay
MSFC/PH membrane (Millipore)

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# Certificate of Analysis

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# **BIOLOGY**

### **Sequence information**

GST-Kinase Recombinant Fusion Protein Amino Acid Sequence							
1					60		
61					120		
121					180		
181					240		
241					300		
301					360		
361					420		
421					480		
481					540		
541					600		
601					660		
1-218: GST	Red: HIS6-tag	Pink: Thrombin cleavage site	Green: 3C/TEV cleavage site	blue: Kinase fragment	boxed		

variation from RefSeq

	Kinase wt <sup>1</sup> Amino Acid Sequence	
1		60
61		120
121		180
181		240
241		300
301		360
361		420
421		480
481		540
541		600
601		660
661		720
721		780
781		840
841		900
901		960
961		1020
1021		1080
1081		1140
1141		1200
1201		1260
1261		1320
1321		1380
1381		1440
1441		1500
1501		1560
1561		1620
	blue: kinase sequence expressed in recombinant protein Red: variant in recombinant protein	

bide. Kindse sequence expressed in recombinant protein . Red. vanant in recomb

<sup>1</sup><u>NCBI/Protein</u> accession number NP\_000000.0 A100B: SNP variation see <u>NCBI/dbSNP</u> ID: rs0000000

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	GST-Kinase Recombinant Fusion Protein Amino Acid Sequence	
1		60
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541		600
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4 040 001	Bad UIO tax Birds Threads a damage of a grant OO/TEV descent of a block (in sec. (a grant of a gran	the second second

### 1-218: GST Red: HIS6-tag Pink: Thrombin cleavage site Green: 3C/TEV cleavage site blue: Kinase fragment box variation from RefSeq

1   60     61   120     121   180     181   240     241   300     301   360     361   420     421   480     481   540     541   600     661   720     721   780     781   840     941   900     961   1020     1021   1080     1081   1140     1241   1320     1321   1380     1321   1380     1321   1320     1501   1500     1501   1500		Kinase wt <sup>2</sup> Amino Acid Sequence	
61   120     121   180     181   240     241   300     301   360     361   420     421   480     481   540     541   600     601   660     661   720     721   780     781   840     900   901     901   960     961   1020     1021   1260     1221   1380     1381   1440     1441   1500     1501   1561	1		60
121   180     181   240     241   300     301   360     361   420     421   480     481   540     541   600     601   600     661   720     721   780     781   840     901   960     961   1020     1021   1020     1021   1260     1221   1320     1321   1380     1381   1440     141   1500     1501   1560	61		120
181   240     241   300     301   360     361   420     421   480     481   540     541   600     601   660     661   720     721   780     781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1261   1320     1381   1440     1441   1500     1501   1561	121		180
241   300     301   360     361   420     421   480     481   540     541   600     601   660     661   720     721   780     781   840     841   900     901   960     961   1020     1021   1200     1201   1200     1201   1260     1321   1380     1381   1440     1441   1500     1501   1561	181		240
301   360     361   420     421   480     481   540     541   600     601   660     661   720     721   780     781   840     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1380     1381   1440     1381   1440     1441   1500     1501   1560	241		300
361   420     421   480     481   540     541   600     601   660     661   720     721   780     781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1201   1260     12201   1320     1321   1380     1381   1440     1441   1500     1501   1560	301		360
421   480     481   540     541   600     601   660     661   720     721   780     781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1201   1260     12201   1320     1321   1380     1381   1440     1441   1500     1501   1560	361		420
481   540     541   600     601   660     661   720     721   780     781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1321   1380     1381   1440     1441   1500     1501   1560	421		480
541   600     601   660     661   720     721   780     781   840     841   900     901   960     961   1020     1021   1081     1141   1140     1201   1200     1221   1320     1321   1380     1381   1440     1441   1500     1501   1560	481		540
601   660     661   720     721   780     781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1321   1380     1381   1440     1441   1500     1501   1560	541		600
661   720     721   780     781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	601		660
721   780     781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1321   1380     1381   1440     1441   1500     1501   1620	661		720
781   840     841   900     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	721		780
841   900     901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1261   1320     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	781		840
901   960     961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1261   1320     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	841		900
961   1020     1021   1080     1081   1140     1141   1200     1201   1260     1261   1320     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	901		960
1021   1080     1081   1140     1141   1200     1201   1260     1261   1320     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	961		1020
1081   1140     1141   1200     1201   1260     1261   1320     1321   1380     1381   1440     1441   1500     1501   1560	1021		1080
1141   1200     1201   1260     1261   1320     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	1081		1140
1201 1260   1261 1320   1321 1380   1381 1440   1441 1500   1501 1560   1561 1620	1141		1200
1261   1320     1321   1380     1381   1440     1441   1500     1501   1560     1561   1620	1201		1260
1321 1380   1381 1440   1441 1500   1501 1560   1561 1620	1261		1320
1381 1440   1441 1500   1501 1560   1561 1620	1321		1380
1441 1500   1501 1560   1561 1620	1381		1440
1501 1561 1561	1441		1500
1561 1620	1501		1560
	1561	blue bisses second a supervision plant sector. Deduced a second is second to be	1620

<sup>2</sup><u>NCBI/Protein</u> accession number NP\_000000.0 A100B: SNP variation see <u>NCBI/dbSNP</u> ID: rs0000000 HGNC identifier #COFACTOR#: YYYY **Recombinant Proteins** 

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..:REACTION