>SOD2_human (NP_000627.2):

MLSRAVCGTSRQLAPVLAYLGSRQKHSLPDLPYDYGALEPHI NAQIMQLHHSKHHAAYVNNLNVTEEKYQEALAKGDVTAQIA LQPALKFNGGGHINHSIFWTNLSPNGGGEPKGELEAIKRDFGS FDKFKEKLTAASVGVQGSGWGWLGFNKERGHLQIAACPNQD PLQGTTGLI

PLLGIDVWEHAYYLQYKNVRPDYLKAIWNVINWENVTERYM ACKK

>SOD2_mouse (NP_038699.2):

MLCRAACSTGRRLGPVAGAAGSRHKHSLPDLPYDYGALEPHI NAQIMQLHHSKHHAAYVNNLNATEEKYHEALAKGDVTTQV ALQPALKFNGGGHINHTIFWTNLSPKGGGEPKGELLEAIKRDF GSFEKFKEKLTAVSVGVQGSGWGWLGFNKEQGRLQIAACSN QDPLQGTTG

LIPLLGIDVWEHAYYLQYKNVRPDYLKAIWNVINWENVTERY TACKK

Score: 1091

Percentage: 10.36%

>SOD2_human (NP_000627.2):

MLSRAVCGTSRQLAPVLAYLGSRQKHSLPDLPYDYGALEPHI NAQIMQLHHSKHHAAYVNNLNVTEEKYQEALAKGDVTAQIA LQPALKFNGGGHINHSIFWTNLSPNGGGEPKGELLEAIKRDFG SFDKFKEKLTAASVGVQGSGWGWLGFNKERGHLQIAACPNQ DPLQGTTGLI

PLLGIDVWEHAYYLQYKNVRPDYLKAIWNVINWENVTERYM ACKK

>RandomSeq:

RHQNRFAPLMEVGCRCMCWFHDWWVISVGTWLHTVIMYM MWPKRFHHNECPKACFRTTYTRKNHHALYWMLFEMCCYDQ DVVWSKTHIFTTVRDIEVYVEQVFFIWGPLCHVAIACYEPVKT

IRRRIPMYLCRHCIRGDNSYLLACCSIIYYFYHHMSYYGVLDIL

Score: -250

Percentage: 94.59%

>SOD2_mouse (NP_038699.2):

MLCRAACSTGRRLGPVAGAAGSRHKHSLPDLPYDYGALEPHI NAQIMQLHHSKHHAAYVNNLNATEEKYHEALAKGDVTTQV ALQPALKFNGGGHINHTIFWTNLSPKGGGEPKGELLEAIKRDF GSFEKFKEKLTAVSVGVQGSGWGWLGFNKEQGRLQIAACSN QDPLQGTTG

LIPLLGIDVWEHAYYLQYKNVRPDYLKAIWNVINWENVTERY TACKK

>RandomSeq:

RHQNRFAPLMEVGCRCMCWFHDWWVISVGTWLHTVIMYM MWPKRFHHNECPKACFRTTYTRKNHHALYWMLFEMCCYDQ DVVWSKTHIFTTVRDIEVYVEQVFFIWGPLCHVAIACYEPVKT IRRRIPMYLCRHCIRGDNSYLLACCSIIYYFYHHMSYYGVLDIL

Score: -250

Percentage: 94.14%