

Product datasheet for TP505984

Bscl2 (NM_008144) Mouse Recombinant Protein

Product data:

Product Type:	Recombinant Proteins
Description:	Purified recombinant protein of Mouse Berardinelli-Seip congenital lipodystrophy 2 (seipin) (Bscl2), with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug
Species:	Mouse
Expression Host:	HEK293T
Expression cDNA Clone or AA Sequence:	>MR205984 representing NM_008144 Red =Cloning site Green =Tags(s)

MIHQRRREAGARETCRDQIKGSDKDEEPSAALSHGQGYRPCGRPARNSKPEAGARPPAVPIMVNDPPVPAL
LWAQEVGHVLAGRARRMLQFGVLFCTILLLLWVSVFLYGSFYYSYMPTVSHLSPVHFHYRTDCDSSTAS
LCSFPVANVSLAKSGRDRVLMYGQPYRVTLLELELPESPVNQDLGMFLTVSCYTRGGRIISTSSRSVMLH
YRSQLLQVLDLTLFSSLLLFGFAEQKQLLEVELYSDYRENSYVPTTGAIIIEHSKRIQMYGAYLRIHAHF
TGLRYLLYNFPMTCAFVGVASNFTFLSVIVLFSYMQVWVWVWPRHRFSLQVNIQRDNNSHHGAPRRISR
HQPGQESTQQSDVTEDGESPEDPSGTEGQLSEEEKPEKRPLNGEEEEQEPEASDGSWEDAALLTEANPPTS
ASASALAPETLGSLRQRPTCSSS

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag:	C-MYC/DDK
Predicted MW:	50 kDa
Concentration:	>0.05 µg/µL as determined by microplate BCA method
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol
Note:	For testing in cell culture applications, please filter before use. Note that you may experience some loss of protein during the filtration process.
Storage:	Store at -80°C after receiving vials.
Stability:	Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.
RefSeq:	NP_032170
Locus ID:	14705



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UniProt ID: [Q9Z2E9](#), [A0A0R4J225](#)

RefSeq Size: 1815

Cytogenetics: 19 5.76 cM

RefSeq ORF: 1329

Synonyms: 2900097C17Rik; AI046355; Gng3lg

Summary: Is a regulator of lipid catabolism essential for adipocyte differentiation. Necessary for correct lipid storage and lipid droplets maintenance; may play a tissue-autonomous role in controlling lipid storage in adipocytes and in preventing ectopic lipid droplet formation in non-adipose tissues. May also be involved in the central regulation of energy homeostasis.
[UniProtKB/Swiss-Prot Function]