double-stranded RNA binding, GO:0003725

RNA phosphodiester bond hydrolysis, GO:0090502 gene silencing by RNA, GO:0031047 core promoter binding, GO:0001047 RNA phosphodiester bond hydrolysis, GO:0090501 response to interferon-alpha, GO:0035455 positive regulation of innate immune response, GO:0045089 protein kinase inhibitor activity, GO:0004860 miRNA metabolic process, GO:0010586 L-alpha-amino acid transmembrane transport, GO:1902475 positive regulation of myelination, GO:0031643 neutral amino acid transmembrane transporter activity, GO:0015175 type B pancreatic cell proliferation, GO:0044342 adenosine deaminase activity, GO:0004000 tRNA dihydrouridine synthase activity, GO:0017150 tRNA dihydrouridine synthesis, GO:0002943 calcium:sodium antiporter activity, GO:0005432 muscle filament sliding, GO:0030049 eye development, GO:0001654