Resource Summary Report

Generated by dkNET on Apr 22, 2025

Hybrigenics

RRID:SCR_003822
Type: Tool

Proper Citation

Hybrigenics (RRID:SCR_003822)

Resource Information

URL: http://www.hybrigenics.com/

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Description: A bio-pharmaceutical company with a focus on research and development of new targets & therapies against proliferative diseases. Its current development program is based on inecalcitol, a vitamin D receptor agonist being studied in three potential indications: * alone in chronic lymphocytic leukaemia * in combination with imatinib in chronic myeloid leukaemia * in prostate cancer, for use with current standards of care Hybrigenics'' research program investigates the action of enzymes called Deubiquitinating Enzymes (DUBs) in the recycling of onco-proteins and the utility of proprietary patented DUB inhibitors against various cancer indications. Hybrigenics Services, a Hybrigenics subsidiary, markets very specialized scientific services to researchers in all areas of life sciences who want to identify, validate and inhibit protein interactions in animal, plant or microbiological cells.

Resource Type: commercial organization

Keywords: enzyme, proteomic, protein interaction network, protein interaction

Related Condition: Lymphocytic leukaemia, Myeloid leukaemia, Prostate cancer, Cancer

Funding:

Resource Name: Hybrigenics

Resource ID: SCR_003822

Alternate IDs: nlx_158127

Record Creation Time: 20220129T080221+0000

Record Last Update: 20250420T014153+0000

Ratings and Alerts

No rating or validation information has been found for Hybrigenics.

No alerts have been found for Hybrigenics.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 15 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Taouis K, et al. (2023) WWOX binds MERIT40 and modulates its function in homologous recombination, implications in breast cancer. Cancer gene therapy, 30(8), 1144.

Hawkins TJ, et al. (2023) NET4 and RabG3 link actin to the tonoplast and facilitate cytoskeletal remodelling during stomatal immunity. Nature communications, 14(1), 5848.

Skeldal S, et al. (2023) Alternative splicing regulates adaptor protein binding, trafficking, and activity of the Vps10p domain receptor SorCS2 in neuronal development. The Journal of biological chemistry, 299(9), 105102.

Derevnina L, et al. (2021) Plant pathogens convergently evolved to counteract redundant nodes of an NLR immune receptor network. PLoS biology, 19(8), e3001136.

van der Wal T, et al. (2020) TMEM98 is a negative regulator of FRAT mediated Wnt/ß-catenin signalling. PloS one, 15(1), e0227435.

Madsen P, et al. (2019) HSPA12A targets the cytoplasmic domain and affects the trafficking of the Amyloid Precursor Protein receptor SorLA. Scientific reports, 9(1), 611.

Skouloudaki K, et al. (2019) The apical protein Apnoia interacts with Crumbs to regulate tracheal growth and inflation. PLoS genetics, 15(1), e1007852.

Chen Y, et al. (2018) Phosphorylation of Arabidopsis SINA2 by CDKG1 affects its ubiquitin ligase activity. BMC plant biology, 18(1), 147.

Miserey-Lenkei S, et al. (2017) Coupling fission and exit of RAB6 vesicles at Golgi hotspots

through kinesin-myosin interactions. Nature communications, 8(1), 1254.

Askarian F, et al. (2016) The interaction between Staphylococcus aureus SdrD and desmoglein 1 is important for adhesion to host cells. Scientific reports, 6, 22134.

Bozickovic O, et al. (2015) Cyclin C interacts with steroid receptor coactivator 2 and upregulates cell cycle genes in MCF-7 cells. Biochimica et biophysica acta, 1853(10 Pt A), 2383.

Steglich B, et al. (2015) The Fun30 chromatin remodeler Fft3 controls nuclear organization and chromatin structure of insulators and subtelomeres in fission yeast. PLoS genetics, 11(3), e1005101.

Zhang B, et al. (2014) PIRIN2 stabilizes cysteine protease XCP2 and increases susceptibility to the vascular pathogen Ralstonia solanacearum in Arabidopsis. The Plant journal : for cell and molecular biology, 79(6), 1009.

Merret R, et al. (2013) XRN4 and LARP1 are required for a heat-triggered mRNA decay pathway involved in plant acclimation and survival during thermal stress. Cell reports, 5(5), 1279.

Bowen AJ, et al. (2010) PAH-domain-specific interactions of the Arabidopsis transcription coregulator SIN3-LIKE1 (SNL1) with telomere-binding protein 1 and ALWAYS EARLY2 Myb-DNA binding factors. Journal of molecular biology, 395(5), 937.