Scopus

Documents

1) Shuli, Z., Linlin, L., Li, G., Yinghu, Z., Nan, S., Haibin, W., Hongyu, X.

Bioinformatics and Computer Simulation Approaches to the Discovery and Analysis of
(2022) *Current Pharmaceutical Biotechnology*, 23 (13), pp. 1541-1555.

2) Pérez-Rubio, G., Ponce-Gallegos, M.A., Domínguez-Mazzocco, B.A., Ponce-Gallegos, J., Ga
Valencia, R.

Role of the host genetic susceptibility to 2009 pandemic influenza a h1n1
(2021) *Viruses*, 13 (2), art. no. 344, .

3) Ge, C., Zhang, W., He, R., Cai, H.

Systematic Identification and Comparative Analysis of Human Cartilage-Derived Self-p
by Ankylosing Spondylitis (AS)-Associated HLA-B*27:05 and Non-AS-associated HLA-
(2020) *International Journal of Peptide Research and Therapeutics*, 26 (1), pp. 513-522.

4) Wu, L.

Chapter 1: Regulatory Considerations for Peptide Therapeutics
(2019) *RSC Drug Discovery Series*, 2019-January (72), pp. 1-30.

5) Mohammadi, M., Falak, R., Emameh, R.Z., Maleki, S.J., Kardar, G.A.

Computational analysis of specific ige epitopes responsible for allergy to fish
(2018) *Current Immunology Reviews*, 14 (2), pp. 130-136.

6) Wu, L.C., Chen, F., Lee, S.L., Raw, A., Yu, L.X.

Building parity between brand and generic peptide products: Regulatory and scientific
synthetic peptides
(2017) *International Journal of Pharmaceutics*, 518 (1-2), pp. 320-334.

7) Schrörs, B., Lübcke, S., Lennerz, V., Fatho, M., Bicker, A., Wölfel, C., Derigs, P., Hankeln, T., Wölfel, T.

HLA class I loss in metachronous metastases prevents continuous T cell recognition c human melanoma model
(2017) *Oncotarget*, 8 (17), pp. 28312-28327.

8) Wang, Y., Zhou, P., Lin, Y., Shu, M., Hu, Y., Xia, Q., Lin, Z.

Quantitative prediction of class i MHC/epitope binding affinity using QSAR modeling d structural information
(2015) *Combinatorial Chemistry and High Throughput Screening*, 18 (1), pp. 75-82.

9) Liu, S., Liu, S., Wang, Y., Liao, Z.

The P2/P2' sites affect the substrate cleavage of TNF-α converting enzyme (TACE)
(2014) *Molecular Immunology*, 62 (1), pp. 122-128.

10) Flower, D.R.

Designing immunogenic peptides
(2013) *Nature Chemical Biology*, 9 (12), pp. 749-753.

11) Fulton, K.M., Twine, S.M.

Immunoproteomics: Current technology and applications
(2013) *Methods in Molecular Biology*, 1061, pp. 21-57.

12) Nirmala, S., Sudandiradoss, C.

Prediction of promiscuous epitopes in the e6 protein of three high risk human papillo approach
(2013) *Asian Pacific Journal of Cancer Prevention*, 14 (7), pp. 4167-4175.

13) Ren, Y., Tian, F., Zhou, P.

Computational peptidology
(2012) *Progress in Chemistry*, 24 (9), pp. 1674-1682.

14) Fadda, L., Körner, C., Kumar, S., van Teijlingen, N.H., Piechocka-Trocha, A., Carrington, M.,

HLA-Cw*0102-restricted HIV-1 p24 epitope variants can modulate the binding of the in and primary NK cell function
(2012) *PLoS Pathogens*, 8 (7), art. no. e1002805, p. 40.

15) Knapp, B., Giczi, V., Ribarics, R., Schreiner, W.

PeptX: Using Genetic Algorithms to optimize peptides for MHC binding
(2011) *BMC Bioinformatics*, 12, art. no. 241, .

16) Li Pira, G., Ivaldi, F., Moretti, P., Manca, F.

High throughput T epitope mapping and vaccine development
(2010) *Journal of Biomedicine and Biotechnology*, 2010, art. no. 325720, .