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**Editorial** 

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### NRF2 molecule of the month

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NRF2 – nuclear factor erythroid 2-related factor 2 is also known as GA binding protein transcription factor, alpha subunit 60kDa- GABPA **[1, 2]**. A recent literature review suggests that NRF2 has complex regulatory roles in cancer. NRF2 is a transcription factor and is upregulated by oncogenes. NRF2 is an antioxidant master regulator and for example, binds to ARE –antioxidant response element. NFR2 lowers ROS (in pancreatic and lung cancers). However, NRF2 silences miR200a (a miRNA) (in breast cancer) that leads to additional expression of its repressor, KEAP1. Therapies are under investigation **[1]**.

 NFE2
 PPARG
 MARE
 PUCIT PAVS
 KAT2A
 MARE
 POL2
 PLS1
 ZUF24
 MAH A2
 DHABI4

 HHOM1
 BATR
 SAT1
 SMARCA4
 FOSL1
 SMAB3
 STATL
 SM55
 DCUMD
 KAP11

 HHOM1
 KEP11
 HEX
 SMARCA4
 FOSL1
 SMAB3
 STATL
 SM55
 DCUMD
 KAP11

 MFE2
 ULUE
 HEX
 SMARCA4
 FOSL1
 SMAB3
 STATL
 SM55
 DCUMD
 KAP11

 MFE2
 ULUE
 HEX
 SMARCA4
 FOSL1
 SMARCA4
 FOSL2
 PRAF
 DCUMD
 KAP11

 MFE2
 ULUE
 HEX
 SMARCA4
 FOSL3
 FOSC
 FOSC4
 FOS

**Figure 1:** NRF2and related interactions with up to 100 neighbors. In this figure, line-colors and various interactions with other genes are red Down-regulation, green Up-regulation, beige Regulation, purple Co-expression, brown Physical ISSN 0973-2063 (online) 0973-8894 (print) Bioinformation 8(18): 846-847 (2012) 8-

Interaction, turquoise dotted Predicted Protein Interaction, and mauve dotted Predicted TFactor Regulation. (GenePro SA Biosciences, http://www.sabiosciences.com/).

The two figures illustrate various gene interactions, up to 100 for each figure. **Figure 1** shows interactions for NRF2 and GABPA that the database treats as similar. **Figure 2** shows interactions and neighboring genes among NRF2-GABPA, KEAP1, NFE2L2, ROS, miR200a, COPD, and MS.



**Figure 2:** NRF2 and related interactionswith NRF2, GABPA, NFE2L2, KEAP1, ROS, miR200a, COPD, MS input) and up to 100 neighbors. In this figure, line-colors and various interactions with other genes are red Down-regulation, green Up-regulation,

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beige Regulation, purple Co-expression, brown Physical Interaction, turquoise dotted Predicted Protein Interaction, and mauve dotted Predicted TFactor Regulation. (GenePro SA Biosciences, http://www.sabiosciences.com/).

It is left as a puzzle for the interested reader to identify the various genes and their functions in the figures [3, 4].

### Acknowledgment:

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### **References:**

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  http://www.genecards.org/cgi-
- bin/carddisp.pl?gene=GABPA&search=NRF2+
- [3] http://www.sabiosciences.com/
- [4] http://www.genecards.org/

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