



1 Article

Structural Insight on Functional Regulation of Human MINERVA Protein

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16 Supplementary Data



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Figure S1. K-M plots of overall survival rate of breast invasive carcinoma (BRCA), glioblastoma
 multiforme (GBM), pancreatic adenocarcinoma (PAAD) patients expressing MINERVA (FAM129B).
 Patients expressing high levels of MINERVA show lower survival rate.



Figure S2. Multiple sequence alignment of FAM129 family members MINERVA (FAM129B), FAM129A, and FAM129C. Regions corresponding to the structured domain of MINERVA (Gly2–Arg560) are well-conserved among the family members. Identical and similar residues are highlighted in red and yellow, respectively. PH domain and the C-terminal flexible region are indicated by blue and orange boxes, respectively. The C-terminal flexible regions vary greatly in length and sequence.

(a)

α2 α8 Arg114 Q302' Q296' Arg86 R306' N137 E133 D129 GIn79 Q262 a Ser82 Glu80 Asp81 β5 α4 α12 **B6** α8 β7 α5

(b)



32 Figure S3. PH domain of MINERVA. (a) Superposition of the PH domains from MINERVA (white 33 cartoon and stick representations) and Pleckstrin (yellow surface representation) in the same view as 34 Figure 3b, left. Side chains of residues that give rise to ligand specificity to MINERVA PH domain are 35 shown in stick representations and labeled (Gln79, Glu80, Asp81, and Ser82). Pleckstrin possesses 36 Gly255, His256, Arg257, and Arg258 on the corresponding positions, that allow larger binding pocket 37 as displayed by the surface representation. (b) Intermolecular interaction between two adjacent 38 molecules of MINERVA establish multiple hydrogen bonds, indicated with yellow dashed lines. The 39 PH domain of a MINERVA molecule is shown as a cartoon representation in cyan-to-green color 40 while helix $\alpha 8'$ of the adjacent MINERVA is in white. The unmodeled region (Ser146–Ile151) are 41 indicated with gray dashed ribbon.

		2019.9.27 Cell 1450-48 BV.		
1, 193 2, Heia 3, Hotiil 4, Asag 5: BxPC3 6, MZA 7. 6965	8. 5F(88 9. 083MG (0. 0333mg 11. MCFN 12. MCGEN 13. ML231 14. 293FT	MINERVA GAPDH		N: FAMI29B
			(a)	





Figure S4. Full Western blots for Figure 1a from same membrane. (a) Long exposure. (b) Medium
 exposure. (c) Short exposure for GAPDH.





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Figure S5. Full Western blot for Figure 1b.









51 Figure S6. Full Western blots for Figure 1d. Sequential blots against (a) anti-PARP, (b) anti-MINERVA
52 (FAM129B), and (c) anti-Actin (loading control).



55 Figure S7. Full Western blots for Figure 5a. Sequential blots on same membrane against (a) anti-FLAG,
56 (b) anti-HA, and (c) anti-Actin (loading control).