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2	Supporting Information
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4	FADD regulates NF- $\kappa$ B activation and promotes ubiquitination of cFLIP <sub>L</sub> to induce
5	apoptosis
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Figure S1. Expression of FADD in HEK 293T cells (a) Immunostaining of FADD in vector transfected cells (endogenous) and pcDNA-FADD transfected cells, post 48 h of incubation, scale bar- 10µm. The pcDNA-FADD was overexpressed for 24-96 h and (b) Percent apoptotic cell death (Annexin-V<sup>+</sup>/PI<sup>+</sup>) (c) and Annexin-V /PI staining were analyzed by Tali cytometer, control represents vector transfected cells. Error bars represent mean±SD, \*P  $\leq 0.05$ , n  $\geq$  3, where n is number of independent experiment.



Figure S2. The uncropped full-length image of western blot results for Figure 1c, e & f. The
chemiluminescent signals from the exposed membrane were recorded on X-ray films.



Figure S3. FADD regulates pro-survival response of TNF-a stimulation. TNF-a (10 ng/ml) 37 stimulation to MCF-7 and 48 h of pcDNA-FADD expressed MCF-7 cells to monitor the (a) 38 Expression of p65 and cFLIP<sub>L</sub> and (b) NF- $\kappa$ B luciferase reporter activity, TNF- $\alpha$  untreated MCF-39 7 cells and 48 h of pcDNA-FADD expressed MCF-7 cells (shown as 0 h time point) were taken 40 as control.. TNF-a (10 ng/ml) stimulation to HCT 116 and 48 h of pcDNA-FADD expressed 41 HCT 116 cells to monitor the (c) Expression of p65 and cFLIP<sub>L</sub> and (d) NF-KB luciferase 42 reporter activity, TNF-a untreated HCT 116 cells and 48 h of pcDNA-FADD expressed HCT 43 44 116 cells (shown as 0 h time point) were taken as control. The uncropped full-length blots are

45 presented in supplementary Figure S4. HEK 293T cells were transfected with; pcDNA-FADD; 46 (column 2), siRNA directed against cFLIP<sub>L</sub> (cFLIP<sub>L</sub><sup>KD</sup>; column 3) and pcDNA-FADD together 47 with siRNA-cFLIP<sub>L</sub> (FADD + cFLIP<sub>L</sub><sup>KD</sup>; column 4). Post 48 h of incubation cells were subjected 48 to monitor (e) Percent cell proliferation, (f) Colony forming assay, (g) Percent crystal violet 49 stained colonies and (h) p65 translocation assay, scale bar- 5µm. The vector and non targeting 50 siRNA transfected cells were taken as a control. Error bars represent mean±SD, \*P ≤0.05, n ≥ 3, 51 where n is number of independent experiment.



Figure S4. The uncropped full-length image of western blot results for Figure S3a & c. The
chemiluminescent signals from the exposed membrane were recorded on X-ray films.

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**Figure S5.** The uncropped full-length image of western blot results for Figure 2a, c & g. The positive control represents  $I\kappa B\alpha$  (inhibitor of NF- $\kappa B$ ) transfected HEK 293T cells, post 48 h of incubation. The chemiluminescent signals from the exposed membrane were recorded on X-ray films.



Figure S6. Effects of domains specific mutation in FADD and cFLIP<sub>L</sub> on cell death. The plasmid constructs of wild type (wt) FADD, mutated FADD (FADD-SLT2; S14G, S16A, S18G, L20A and T21V), death domain (DD) of FADD (FADD without DED), wt cFLIP<sub>L</sub>, mutated MC159 (FLIP-DM; an analogous of cFLIP<sub>L</sub> unable to activate NF-κB signaling) were transfected to HEK 293T cells for 48 h and (a) Percent apoptotic death and, (b) Annexin-V-alexa 488/PI staining, detected by Tali cytometer, Error bars represent mean ± SD, \*P ≤0.05, n ≥ 3, where n is number of independent experiment.



(C)

Figure S7. The uncropped full-length image of western blot results for Figure 3c. Thechemiluminescent signals from the exposed membrane were recorded on X-ray films.



Figure S8. Over expression of FADD commences cell death, independent of TNF-a 73 stimulation. HEK 293T cells were treated with TNF- $\alpha$  (5, 10 and 15 ng/ml) for 6-24 h. Cell 74 death was examined by Annexin V /PI staining detection by automated cell counter Tali<sup>TM</sup> (a) 6 75 h, (b) 12 h, (c) 24 h, (d) Percent cell death by by automated cell counter Tali<sup>TM</sup> and (f) TNF- $\alpha$ 76 (10 ng/ml) stimulation to 48 h of pcDNA-FADD transfected cells to monitor the cell viability in 77 (a) MCF-7, (g) HCT 116 cells and compared with TNF- $\alpha$  treated non-transfected cells. TNF- $\alpha$ 78 untreated and 48 h of FADD expressed cells (shown as 0 h time) were taken as controls. Error 79 bars represent mean $\pm$ SD, \*P  $\leq 0.05$ , n  $\geq 3$ , where n is number of independent experiment. 80



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Figure S9. Ablation of NF- $\kappa$ B activation and cFLIP<sub>L</sub> expression commences apoptotic cell death. TNF- $\alpha$  (10 ng/ml) stimulation to HEK 293T cells and 48 h of pcDNA-FADD expressed HEK 293T cells. (a) Colony forming assay, (b) Percent crystal violet stained colonies (c) FACS analysis of individually TNF- $\alpha$  treated and FADD expressed cells by BD FACSCalibur<sup>TM</sup>, (d)

86	Annexin-V-FITC/PI staining detection by Tali cytometer, (e) The pEGFP-cytochrome c stable
87	HEK 293T cells were transfected with pcDNA-FADD (48 h) and further stimulated with TNF- $\alpha$
88	(10 ng/ml) (f) HEK 293T cells were transfected with pEYFP-FADD (48 h) and treated with
89	TNF- $\alpha$ (10 ng/ml), post incubation cells were stained with MitoTracker Red and counterstained
90	with DAPI, the cells were analyzed under fluorescent microscope, control represents vector
91	transfected cells (shown as 0 min time), scale bar- 5µm. HEK 293T cells were treated with,
92	TNF- $\alpha$ (10 ng/ml) for 12 h ( <i>lane 2</i> ), siRNA-cFLIP <sub>L</sub> (cFLIP <sub>L</sub> <sup>KD</sup> ; <i>lane 3</i> ) for 48 h and TNF- $\alpha$ (10
93	ng/ml) primed (12 h) cells followed by siRNA-cFLIP <sub>L</sub> (TNF- $\alpha$ + cFLIP <sub>L</sub> <sup>KD</sup> ; <i>lane 4</i> ) transfection
94	for an additional 48 h, post incubation, (g) Annexin-V-FITC/PI staining detection by Tali
95	cytometer (h) Qualitative analysis of mitochondrial integrity on pEGFP-cytochrome c stable
96	cells, (i) JC-1 staining. Error bars represent mean±SD, *P $\leq 0.05$ , n $\geq$ 3, where n is number of
97	independent experiment.

![](_page_11_Figure_0.jpeg)

Figure S10. The uncropped full-length image of western blot results for Figure 4f & l. Thechemiluminescent signals from the exposed membrane were recorded on X-ray films.

![](_page_12_Figure_0.jpeg)

**Figure S11. FADD suppresses necrotic death.** HEK 293T cells were pretreated with Necrostatin-1 (20 $\mu$ M) for 18 h and further stimulated with TNF- $\alpha$  (10 ng/ml) in HEK 293T and 48 h of pcDNA-FADD expressed HEK 293T cells to monitor (a) Percent LDH release and, (b) Cells were lyzed and the activation of PARP was monitored by western blot analysis. Control represents untreated and FADD expressed cells for 48 h (shown as 0 min time). Error bars represent mean±SD, \*P ≤0.05, n ≥ 3, where n is number of independent experiment.

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![](_page_13_Figure_0.jpeg)

Figure S12. FADD regulates expression of apoptotic components. TNF-a (10 ng/ml) 117 stimulation to HEK 293T and 48 h of pcDNA-FADD expressed HEK 293T cells to analyze the 118 119 (a) mRNA expression of cFLIP<sub>L</sub> (b) mRNA expression of FADD (c) mRNA expression of RIP1 and (d) mRNA expression of Procaspase-8. The values were normalized by using the difference 120 in critical thresholds (C<sub>T</sub>) between target gene and 18SrRNA (endogenous control). The 121 expression of mRNA of the respective genes was compared with untreated and 48 h of FADD 122 expressed cells (shown as 0 min time) control using the values of  $2^{-\Delta\Delta CT}$ . Error bars represent 123 mean $\pm$ SD, \*P  $\leq 0.05$ , \*\*P  $\leq 0.001$ , n  $\geq 3$ , where n is number of independent experiment. 124

![](_page_14_Figure_0.jpeg)

Figure S13. The uncropped full-length image of western blot results for Figure 5g & i. Thechemiluminescent signals from the exposed membrane were recorded on X-ray films.

![](_page_15_Figure_0.jpeg)

**Figure S14. NAC pretreatment restore ROS mediated JNK activation.** HEK 293T cells were pretreated with NAC (25µM) for 3h and further stimulated with TNF- $\alpha$  (10 ng/ml) for 12 h (*bar* 2), transfected with siRNA-cFLIP<sub>L</sub> (cFLIP<sub>L</sub><sup>KD</sup>; *bar* 3) for 48 h and TNF- $\alpha$  (10 ng/ml) primed (12 h) cells transfected with siRNA-cFLIP<sub>L</sub> (TNF- $\alpha$  + cFLIP<sub>L</sub><sup>KD</sup>; *bar* 4) for an additional 48 h. Post incubation the (a) ROS measurement and (b) Western blot analysis of JNK1 and ITCH. The uncropped full-length blot of JNK1 is presented in supplementary Figure S15. Error bars represent mean±SD, \*P ≤0.05, n ≥ 3, where n is number of independent experiment.

![](_page_16_Figure_0.jpeg)

(Figure S13b)

Figure S15. The uncropped full-length image of western blot results for Figure 6h & j and
supplementary fig S13b. The positive control represents cycloheximide (5µg/ml for 8 h) treated
HEK 293T cells. The chemiluminescent signals from the exposed membrane were recorded on
X-ray films.