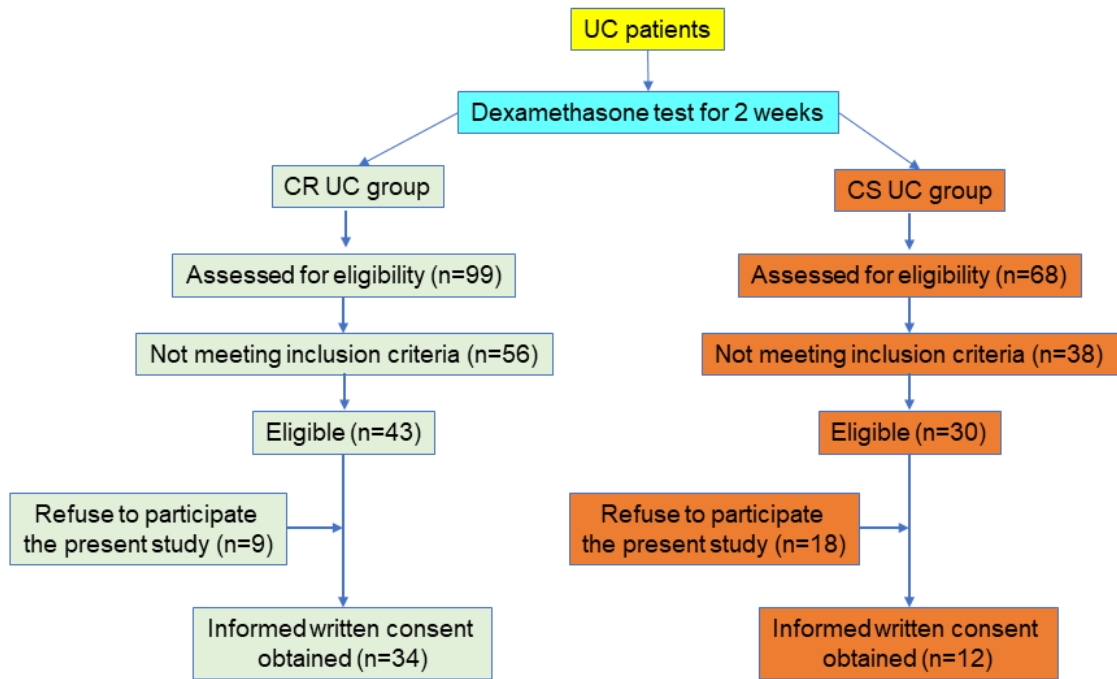**Supplementary materials**

Figure S1. A flow chart shows UC patient selection.
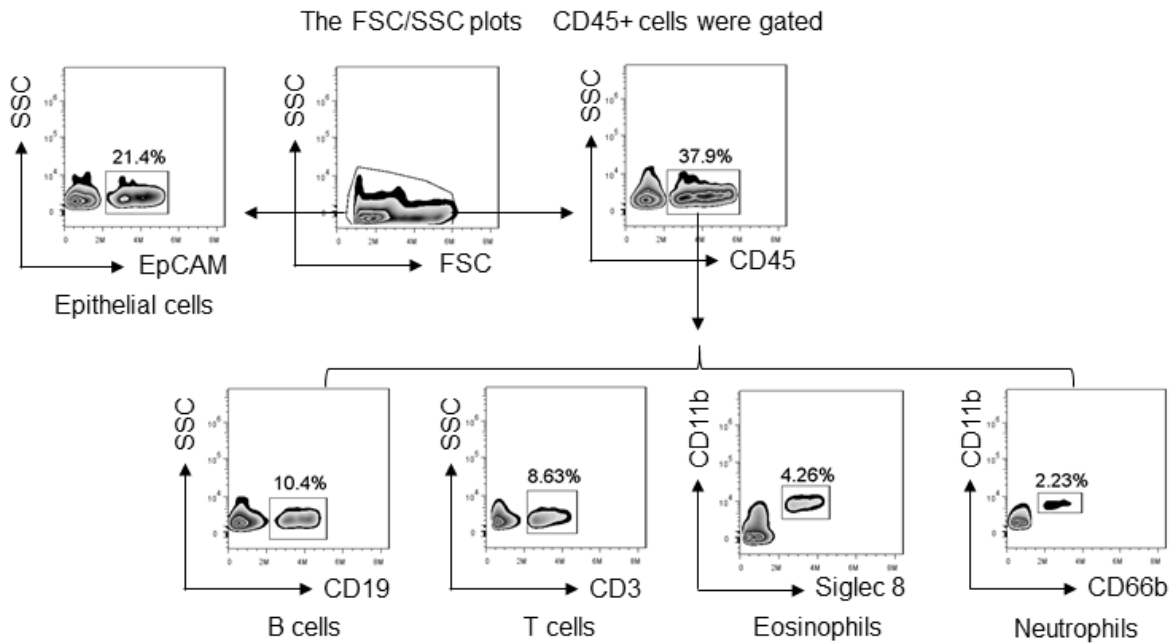

Figure S2. FACS gating procedures. Surgically removed colon tissues from a non-UC patient were used in this experiment to illustrate the FACS gating procedures.
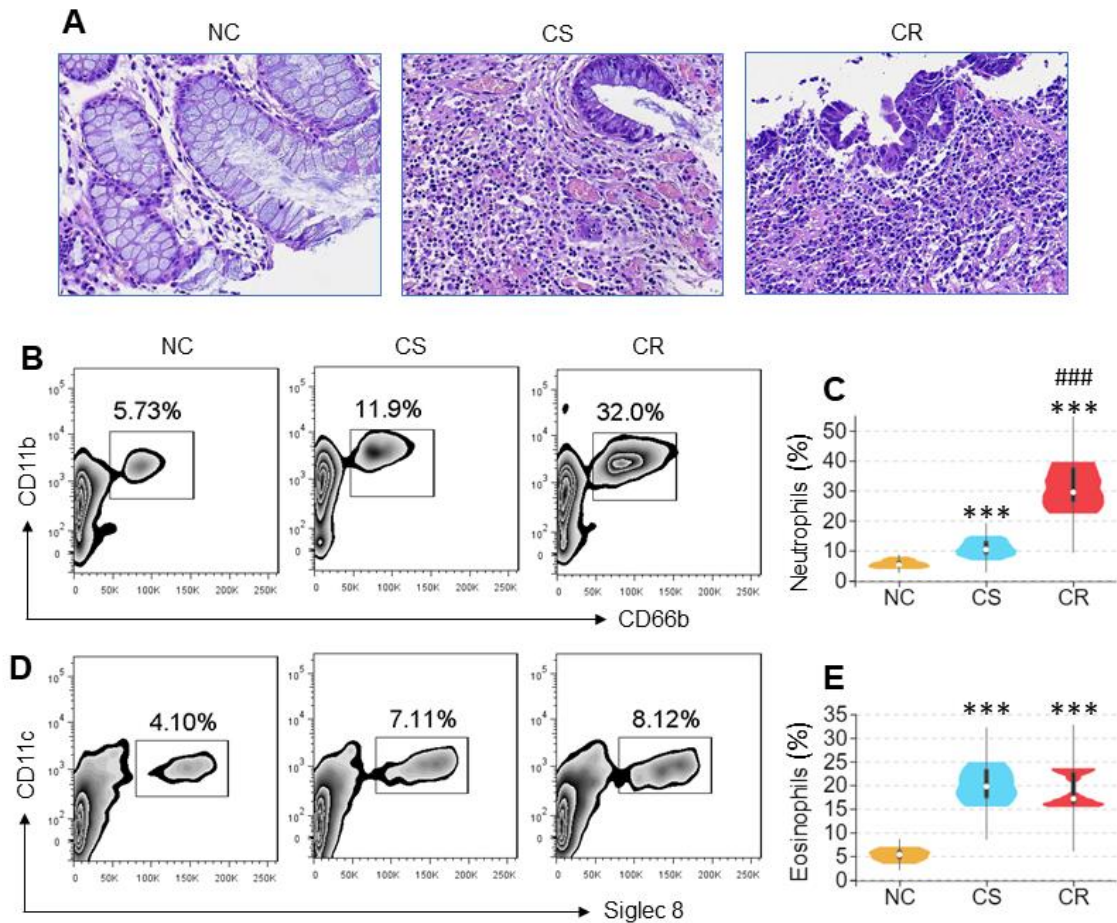
Figure S3. The UC colon tissues of CR patients show high frequency of neutrophils (N). Surgically removed colon tissues were collected from patients with UC and colon cancer, including subjects with corticosteroid resistance (CR group; n=34), or corticosteroid sensitiveness (CS group; n=12). Colon samples from patients with colon cancer without UC were also collected used as non-UC controls (NC; n=8). A, representative colon histology images (original magnification: ×200) show profound inflammatory cell infiltration in the tissues. B-E, lamina propria mononuclear cells (LPMC) were isolated from the colon samples and analyzed by FACS. The gated FACS plots show cell counts of neutrophils (B) and Eo (D). The violin plots show summarized cell counts data of N (C) and Eo (E). ***, p<0.001 (Mann Whitney Test) compared with NC group. ###, p<0.001, compared with CS group. The data of violin plots are presented as mean (IGR) and data range.

Human TW1 amino acid sequence (NM_000474.4)
MMQDVSSS**PVSP**ADDSLSNSEEEPDRQQPPSGKRGGRKRRSSRRSAGGGAGPGGA
AGGGVGGGDE**PGSP**AQGKRGKKSAGCGGGGGGAGGGGGSSSGGGSPQSYEELQTQ
RVMANVRERQRTQSLNEAFAALRKII**PTLP**SDKLSKIQTLKLAARYIDFLYQVLQSDELD
SKMASCSYVAHERLSYAFSVWRMEGAWSMSASH

Figure S4. Three PxxP structures in the amino acid sequence of TW1. Three PxxP structures (in red) in the amino acid sequence of TW1 molecule.

Peptide 1   Peptide 2

NLTSLGTLNFPGRTVFSN   FPQQGQISLSSGETDLK

Human GRα molecular amino acid sequence (GenBank: U01351.1)
MDSKESLTPGREENPSSVLAQERGDVMDFYKTLRGGATVKVSASSPSLAVASQSDSK
QRRLLVDFPKGSVSNAQQPDLSKAVSLSMGLYMGETETKVMGNDLG**FPQQGQISLSS
GETDLK**LLEESIANLNRSTSVPENPKSSASTAVSAAPTEKEFPKTHSDVSSEQQHLKGQ
TGTNGGNVKLYTTDQSTFDILQDLEFSSGSPGKETNESPWRSDLL
IDENCLLSPLAGEDDSFLLEGNSNEDCKPLILPDTKPKIKDNGDLVLSSPSNVTLPQVKT
EKEDFIELCTPGVIKQEKLGTVYCQASFPGANIIGNKMSAISVHGVSTSGGQMYHYDMN
TASLSQQQDQKPIFNVIPPIPVGSENWNRCQGSGDD**NLTSLGTLNFPGRTVFSN**GYSS
PSMRPDVSSPPSSSSTATTGPPPKLCLVCSDEASGCHYGVLTCGSCKVFFKRAVEGR
QHNYLCAGRNDCIIDKIRRKNCPACRYRKCLQAGMNLEARKTKKKIKGIQQATTGVSQE
TSENPGNKTIVPATLPQLTPTLVSLLEVIEPEVLYAGYDSSVPDSTWRIMTTLNMLGGRQ
VIAAVKWAKAIPGFRNLHLDDQMTLLQYSWMFLMAFALGWRSYRQSSANLLCFAPDLII
NEQRMTLPCMYDQCKHMLYVSSELHRLQVSYEEYLCMKTLLLLSSVPKDGLKSQELF
DEIRMTYIKELGKAIVKREGNSSQNWQRFYQLTKLLDSMHEVVENLLNYCFQTFLDKTM
SIEFPEMLAEIITNQIPKYSNGNIKKLLFHQK

Figure S5. Identification of GRα and in a complex precipitated by an anti-TW1 Ab. A complex in protein extracts of neutrophils isolated from UC colon tissues of patients with CR was analyzed by MS. The two representative peptides indicate that they belong to GRα molecule.
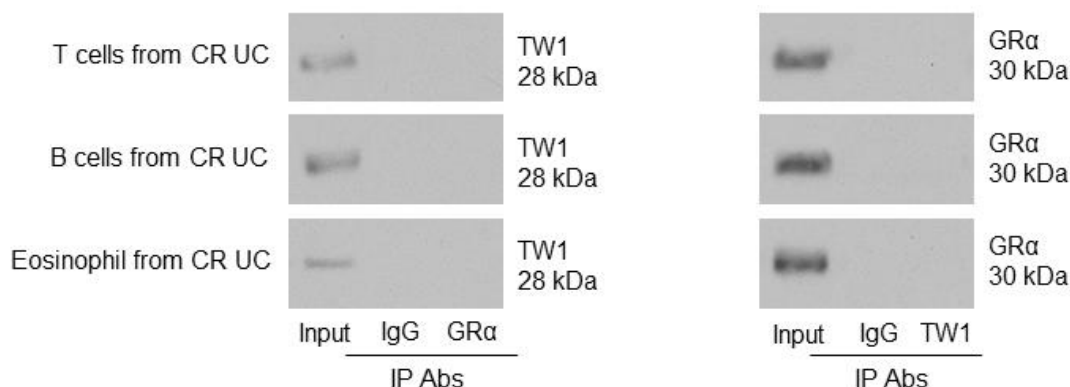
Figure S6. Assessment of TW1/GRα complex in immune cells isolated from CR UC colon tissues. T cells, B cells and eosinophils were isolated from CR UC colon tissues as described and analyzed by co-IP with anti-TW1 Ab and GRα Ab as precipitating Abs and blotting Abs, respectively. Immunoblots show co-IP results. The data are from one experiment that represent 6 independent experiments.
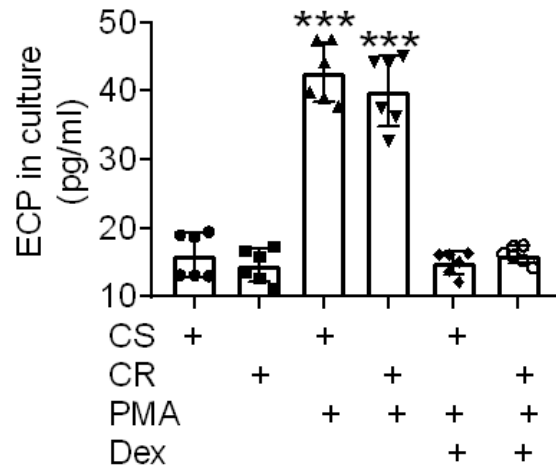

Figure S7. Eosinophils isolated from CR and CS colon tissues do not show CR. Isolated colon eosinophils were treated with the agents listed below the bar graph. CS: Corticosteroid sensitive UC patients. CR: Corticosteroid resistant UC patients. PMA: PMA in the culture at 50 nM. Dex: Dexamethasone in the culture at 1 μM. The bars show ECP levels in the culture. ***, p<0.001 (ANOVA followed by Dunnectt's test), compared with CS alone group.
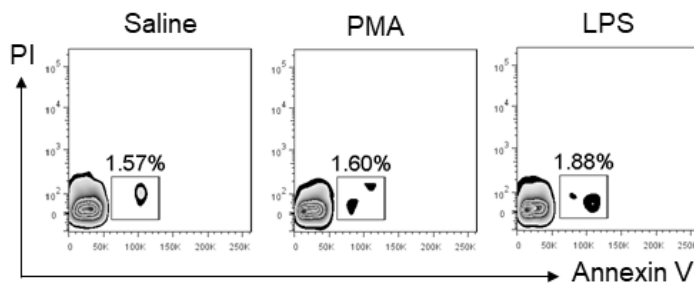

Figure S8. Assessment of neutrophil apoptosis. Isolated CR neutrophils were cultured in the presence of PMA (50 ng/ml) or LPS (250 ng/ml) overnight. The cells were stained with Annexin V reagent and PI, and analyzed by FACS. The gated FACS plots show apoptotic cells. The data represent 3 independent experiments.
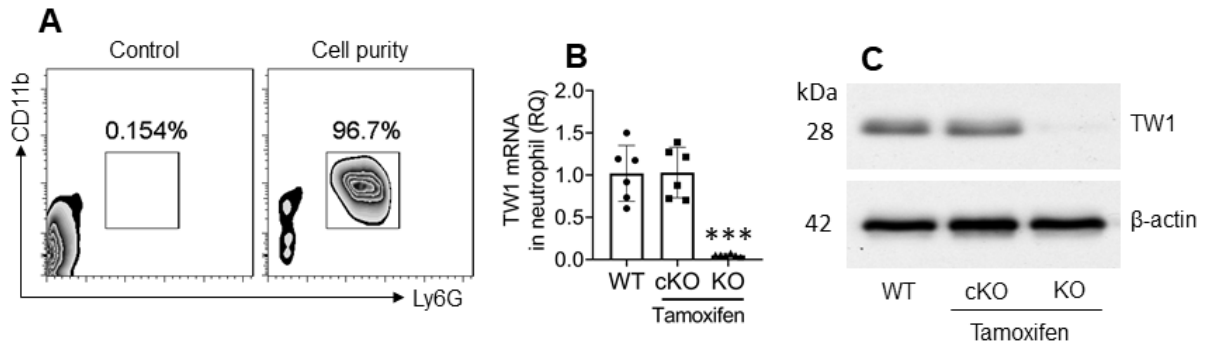
Figure S9. Checking the TW1 expression in neutrophils of mice carrying the TW1-knockout neutrophils. Neutrophils were isolated from the colon tissues of WT mice, mice carrying neutrophils have a WT TW1 gene and a Cre gene (cKO), and mice carrying the TW1-knockout neutrophils (KO) by FACS. A, the gated plots show cell purity of isolated neutrophils. B, the bars show TW1 mRNA levels in neutrophils. C, the immunoblots show TW1 protein levels in neutrophils. Tamoxifen: Mice were treated with tamoxifen daily for one week to activate the TW1-gene-depletion.

Peptide 1
CPPAMRLAFKQLHRRV

Peptide 2
VFWGSQSLETSTIKKT

RASAL1 amino acid sequence (GenBank: BC093724.1)
MAKSSSLNVRVVEGRALPAKDVSGSSDPYCLVKVDDEVVARTATVWRSLGPFWGEEY
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GEICLSVQMLEDGQGRCLRCHVLQARDLAPRDISGTSDPFARVFWGSQSLETSTIKKT
RFPHWDEVLELREMPGAPSPLRVELWDWDMVGKNDFLGMVEFSPKTLQQKPPKGW
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MAPLHPFLLQCVSRVRDFLDRLVDVDGDEAGVPARALFPPSAIVREGYLLKRKEEPAG
LATRFAFKKRYVWLSGETLSFSKSPEWQMCHSIPVSHIRAVERVDEGAFQLPHVMQV
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LQAERSAAGCSRTHSAVTLGDWSDPLDPDAEAQTVYRQLLLGRDQLRLKLLEDSNMD
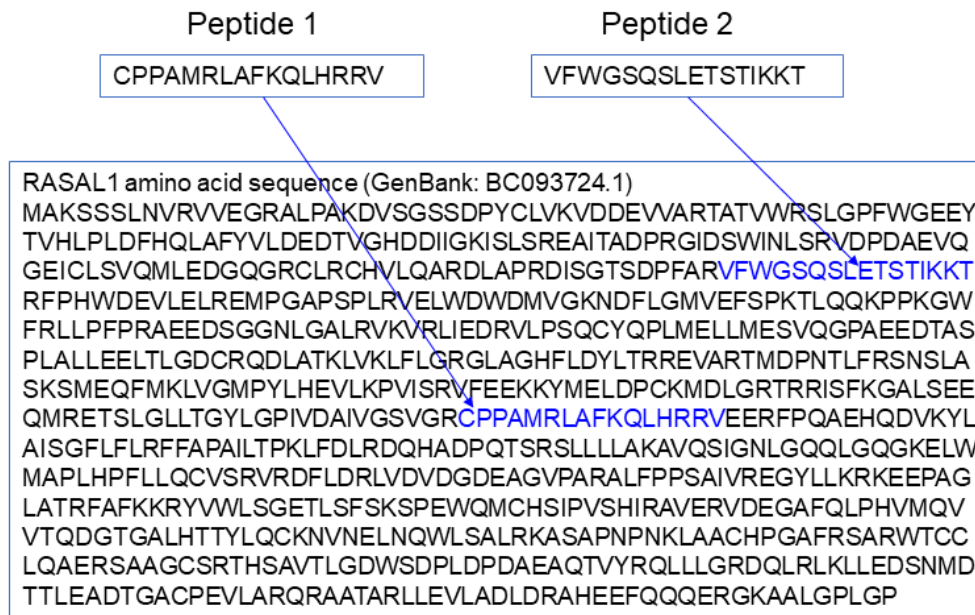TTLEADTGACPEVLARQRAATARLLEVLADLDRAHEEFQQQERGKAALGPLGP

Figure S10. Identification of proteins bound to STAT3 in neutrophils. Proteins were extracted from neutrophils isolated from the colon tissues of UC patients with CR and analyzed by MS. The two representative peptides (the arrows point the original sites; in blue) indicate the protein bound STAT3 is RASAL1L.

Human STAT3) amino acid sequence (NM_139276.3)
MAQWNQLQQLDTRYLEQLHQLYSDSFPMELRQFLAPWIESQDWAYAASKESHATLVFHNLL
GEIDQQYSRFLQESNVLYQHNLRRIKQFLQSRYLEKPMEIARIVARCLWEESRLLQTAATAAQ
QGGQANHPTAAVVTEKQQMLEQHLQDVRKRVQDLEQKMKVVENLQDDFDFNYKTLKSQG
DMQDLNGNNQSVTRQKMQQLEQMLTALDQMRRSIVSELAGLLSAMEYVQKTLTDEELADW
KRRQQIACIGGPPNICLDRLENWITSLAESQLQTRQQIKKLEELQQKVSYKGDPIVQHRPMLE
ERIVELFRNLMKSAFVVERQ**PCMPMHP**DRPLVIKTGVQFTTKVRLLVKFPELNYQLKIKVCID
KDSGDVAALRGSRKFNILGTNTKVMNMEESNNGSLSAEFKHLTLREQRCGNGGRANCDAS
LIVTEELHLITFETEVYHQGLKIDLETHSLPVVVISNICQMPNAWASILWYNMLTNNPKNVNFFT
KPPIGTWDQVAEVLSWQFSSTTKRGLSIEQLTTLAEKLLGPGVNYSGCQITWAKFCKENMAG
KGFSFWVWLDNIIDLVKKYILALWNEGYIMGFISKERERAILSTKPPGTFLLRFSESSKEGGVT
FTWVEKDISGKTQIQSVEPYTKQQLNNMSFAEIIMGYKIMDATNILVSPLVYLY**PDIP**KEEAFGK
YCRPESQEHPEADPGSAAPYLKTKFICVTPTTCSNTIDL**PMSP**RTLDSLMQFGNNGEGAEPS
AGGQFESLTFDMELTSECATSPM

Figure S11. PxxP structures in STAT3 amino acid sequence. There are four "PxxP" structures (in red) in the STAT3 amino acid sequence.
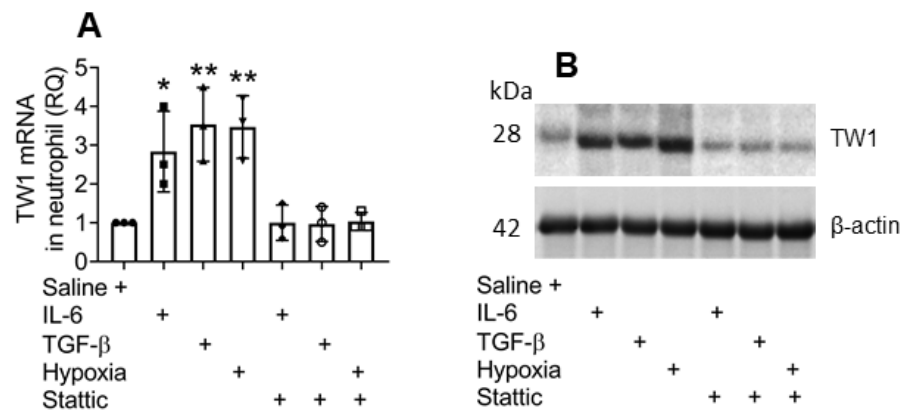
Figure S12. Inhibition of STAT3 attenuates TW1 expression in neutrophils. Neutrophils were isolated from blood samples obtained from healthy human subjects and exposed to IL-6 (50 ng/ml) or TGF-β (10 ng/ml) in the culture, or treated with hypoxia for 24 h. Stattic (an inhibitor of STAT3; 1 μM) was added to the culture in separate wells of each experimental setting. A, the bars indicate TW1 mRNA levels in neutrophils. B, the immunoblots indicate TW1 protein levels in neutrophils. *, p<0.01 (ANOVA followed by the Dunnett's test), compared with the saline group.

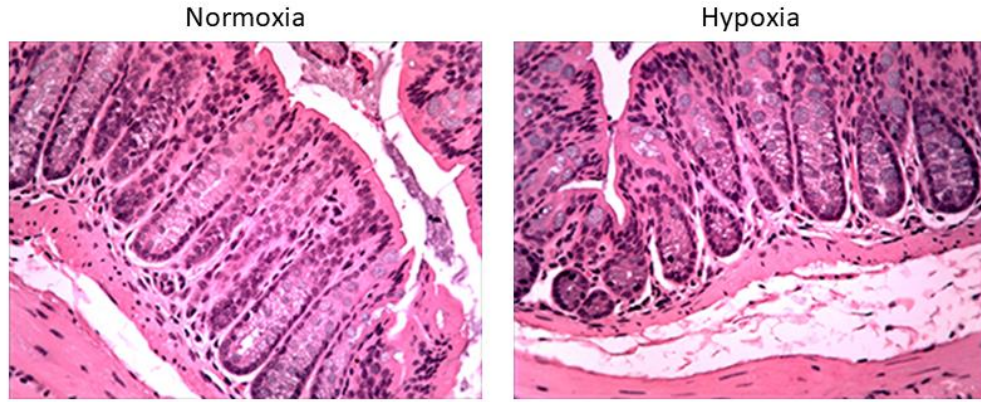Figure S13. Colon histology of mice treated with normoxia or hypoxia. Used as controls, naïve BALB/c mice were treated with normoxia alone or hypoxia alone daily for one week. Colon segments were excised and processed for paraffin section and HE staining. The representative images show colon tissue structure. Original magnification: ×200.
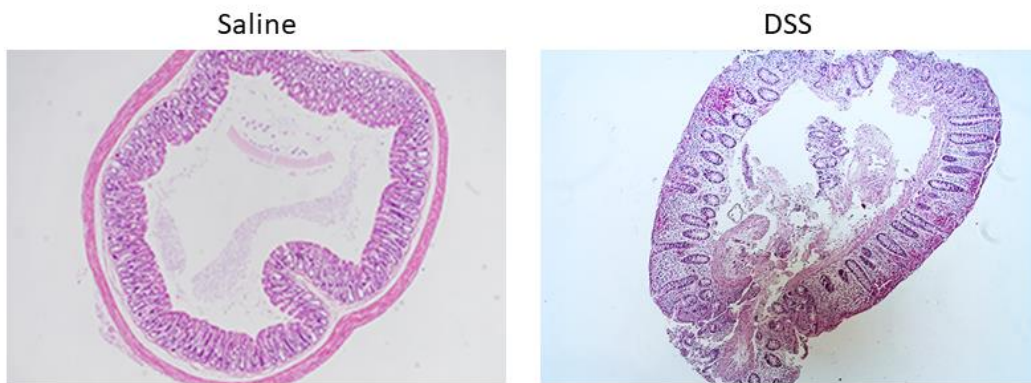
Figure S14. Colon histology of mice carrying TW1-deficient neutrophils. Mice carrying TW1-deficient neutrophils were treated with saline or DSS in the same procedures as described in Fig. 6. Representative images show histology of the colon. Original magnification: ×50. Each group consists of 6 mice.
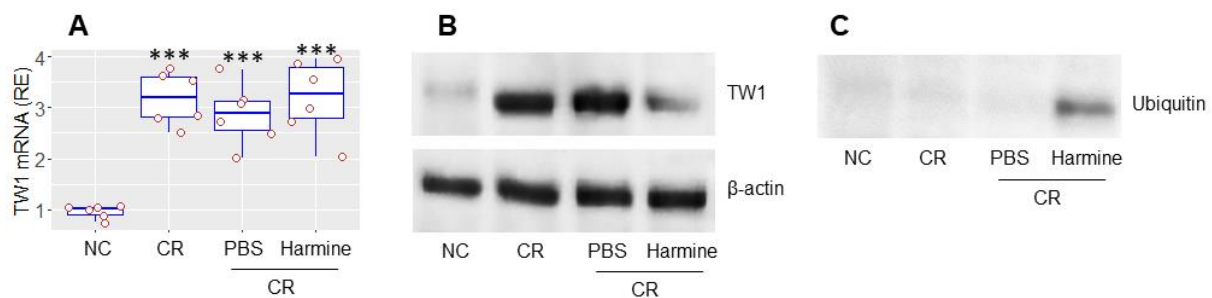
Figure S15. Assessment of the role of harmine in regulating TW1 in neutrophils. Neutrophils were isolated from surgically removed colon tissues of NC (n=6) subjects and CR/UC subjects (n=6), and cultured with the conditions denoted below the graphs for 48 h. Neutrophils were

harvested at the end of culture and analyzed by RT-qPCR and Western blotting. A, boxplots show TW1 mRNA levels in neutrophils. B, immunoblots show TW1 protein levels in neutrophils. C, the PVDF membrane of TW1 blots was processed with the "peel-re-blotting" procedures, and re-blotted with anti-ubiquitin Ab. Immunoblots show ubiquitin levels. Each bubble in boxplots presents data obtained from one sample. ***, p<0.001 (ANOVA + Dunnett's test), compared with the NC group. The data of panel B and C are from one experiment that represent 6 independent experiments. Harmine = 20 μM.

RNAseq data

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2    TLR4  -0.68712598
3    TWIST1  -0.44358750
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11   TNF  0.11251925
12   TLR5  3.12821995
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107   TNF 0.68460297
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116 TLR2 2.62525250
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119 ELANE -0.886840609
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123   TNF 0.36978208
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125   FAS 0.18507487
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133 TLR7 -0.15071077
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349 FAS -0.34877614
350 FASL -0.87878743

351   P53   0.54277769
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353 Case   CR23
354 TLR4   1.01344842
355 TWIST1 -0.96108296
356 TLR2   0.91728069
357 TLR7 -0.97753115
358 NFKB1   2.90912036
359 ELANE 2.359157233
360   MPO   1.49737065
361   GCR   3.27190154
362 KRAS   1.56634159
363   TNF   0.25528179
364 TLR5   1.71635660
365   FAS   0.92514166
366 FASL   0.54578311
367   P53   0.39042885
368 IKBKB -0.678267577
369 Case   CR24
370 TLR4   0.22905628
371 TWIST1 0.74586539
372 TLR2 -0.52328368
373 TLR7   1.61535234
374 NFKB1   0.77452040
375 ELANE 0.584378020
376   MPO   0.16812264
377   GCR   0.19068047
378 KRAS   1.81062305
379   TNF -0.53232207
380 TLR5 -0.99925450
381   FAS   0.09013685
382 FASL   0.92726904
383   P53 -1.48113647
384 IKBKB 0.505167076
385 Case   CR25
386 TLR4   1.21431105
387 TWIST1 2.62719229
388 TLR2   2.17336304
389 TLR7   0.09079613
390 NFKB1 -0.21090343
391 ELANE 2.825430063
392   MPO   0.41060157
393   GCR   3.33952631
394 KRAS   0.98556896

395 TNF 0.61683418
396 TLR5 -0.61492445
397 FAS -1.99764842
398 FASL -0.15718572
399 P53 0.07285287
400 IKBKB -2.061456747
401 Case CR26
402 TLR4 0.14698725
403 TWIST1 -0.55301936
404 TLR2 -0.81125540
405 TLR7 2.49925679
406 NFKB1 3.34156790
407 ELANE 0.751254260
408 MPO 0.20090513
409 GCR 0.03416019
410 KRAS 2.03486147
411 TNF -0.04502768
412 TLR5 -0.17351294
413 FAS 0.18381280
414 FASL -1.35408317
415 P53 -1.91352347
416 IKBKB -1.273856069
417 Case CR27
418 TLR4 2.36372914
419 TWIST1 2.75113189
420 TLR2 2.75087641
421 TLR7 3.48379017
422 NFKB1 -0.24774763
423 ELANE 1.820368859
424 MPO -0.36723892
425 GCR 0.91024786
426 KRAS -0.15652828
427 TNF 2.73206257
428 TLR5 0.65451321
429 FAS -2.15891621
430 FASL -0.75755730
431 P53 -2.03065845
432 IKBKB -0.815185238
433 Case CR28
434 TLR4 -0.69866310
435 TWIST1 1.28120713
436 TLR2 2.35093401
437 TLR7 -0.86847257
438 NFKB1 0.75600575

439 ELANE 3.468022713
440   MPO -0.65587403
441   GCR 1.79955697
442 KRAS 1.82731242
443   TNF 0.25184842
444 TLR5 0.63394880
445   FAS -0.86264272
446 FASL 0.05100667
447   P53 0.77018233
448 IKBKB -1.716161011
449 Case   CR29
450 TLR4 1.32615615
451 TWIST1 -0.57993896
452 TLR2 0.21816052
453 TLR7 -0.20210118
454 NFKB1 1.93586087
455 ELANE 1.704619285
456   MPO 0.07516186
457   GCR 3.26560644
458 KRAS 2.46530833
459   TNF -0.37302067
460 TLR5 -0.71961459
461   FAS -1.43875510
462 FASL -1.29074468
463   P53 -0.26902073
464 IKBKB -1.377418032
465 Case   CR30
466 TLR4 2.01936452
467 TWIST1 -0.52343897
468 TLR2 3.02104546
469 TLR7 2.21026226
470 NFKB1 -0.05774033
471 ELANE 3.193286139
472   MPO 2.90757481
473   GCR 2.43210202
474 KRAS 2.47010841
475   TNF -0.35343804
476 TLR5 1.72056378
477   FAS 0.99992923
478 FASL 0.37878184
479   P53 -0.30977395
480 IKBKB -0.694693767
481 Case   CR31
482 TLR4 -0.20058494

483 TWIST1 2.32623775
484 TLR2 2.92420292
485 TLR7 2.29834789
486 NFKB1 1.28386083
487 ELANE 2.588147438
488   MPO 3.03632227
489   GCR -0.15308277
490 KRAS -0.11909661
491   TNF 1.68656603
492 TLR5 -0.67223278
493   FAS 0.66315863
494 FASL 0.84639040
495   P53 0.46782031
496 IKBKB -0.380537026
497 Case   CR32
498 TLR4 2.07161684
499 TWIST1 0.49027457
500 TLR2 2.08540385
501 TLR7 -0.07046730
502 NFKB1 2.94993429
503 ELANE 2.313198432
504   MPO 1.83010650
505   GCR 3.32159981
506 KRAS 0.14123436
507   TNF -0.56153306
508 TLR5 -0.10509159
509   FAS -0.16160331
510 FASL 0.24487897
511   P53 0.90781418
512 IKBKB -0.338627279
513 Case   CR33
514 TLR4 1.88541973
515 TWIST1 -0.52814760
516 TLR2 0.48060327
517 TLR7 3.31257107
518 NFKB1 1.20630885
519 ELANE -0.835479638
520   MPO 2.62650044
521   GCR -0.01936750
522 KRAS -0.30032743
523   TNF 0.96100580
524 TLR5 0.15230126
525   FAS 0.84738177
526 FASL -1.26953621

527   P53 -0.62598614
528 IKBKB -1.808931093
529 Case   CR34
530 TLR4 1.35451742
531 TWIST1 2.41818137
532 TLR2 1.77238071
533 TLR7 1.06822339
534 NFKB1 -0.29008608
535 ELANE 1.894641702
536   MPO 3.37915944
537   GCR 0.92857771
538 KRAS 1.95369571
539   TNF 0.76152267
540 TLR5 -0.44486944
541   FAS -0.23780702
542 FASL -0.78588475
543   P53 0.54165449
544 IKBKB 0.590722483
545 Case   CS1
546 TLR4 -1.91617819
547 TWIST1 0.21703582
548 TLR2 0.63208493
549 TLR7 -1.05759056
550 NFKB1 -1.60600234
551 ELANE -1.670798733
552   MPO 0.44582402
553   GCR -1.19730863
554 KRAS -0.11408166
555   TNF -1.53079166
556 TLR5 0.11081331
557   FAS 0.01610904
558 FASL 0.04612262
559   P53 0.52558018
560 IKBKB 0.781607668
561 Case   CS2
562 TLR4 0.79064137
563 TWIST1 -0.98791178
564 TLR2 -0.36810504
565 TLR7 -0.87750903
566 NFKB1 -0.07458511
567 ELANE -1.829979494
568   MPO -1.43190058
569   GCR -0.85537168
570 KRAS 0.18653482

571   TNF -0.69888010
572   TLR5 -0.65677541
573   FAS 0.06864689
574   FASL 0.03285200
575   P53 0.52769821
576   IKBKB -0.947274562
577   Case   CS3
578   TLR4 -1.17980306
579   TWIST1 -0.47191294
580   TLR2 0.35140712
581   TLR7 -1.39007637
582   NFKB1 0.77414820
583   ELANE -0.420394886
584   MPO -1.96732515
585   GCR -1.46747333
586   KRAS -0.55083493
587   TNF 0.23908286
588   TLR5 -0.03525878
589   FAS -0.62597222
590   FASL -0.75150608
591   P53 0.05114987
592   IKBKB -0.364605348
593   Case   CS4
594   TLR4 -1.20747959
595   TWIST1 -1.87229378
596   TLR2 -1.11502018
597   TLR7 0.13542458
598   NFKB1 -1.59942666
599   ELANE -1.621675825
600   MPO -1.43304906
601   GCR -1.87418095
602   KRAS -1.75279755
603   TNF 0.80140364
604   TLR5 -0.43072939
605   FAS 0.22156247
606   FASL 0.63431364
607   P53 1.02573213
608   IKBKB -0.106936762
609   Case   CS5
610   TLR4 -0.80678737
611   TWIST1 -0.91524222
612   TLR2 0.83432388
613   TLR7 -1.61513445
614   NFKB1 -0.98566762

615 ELANE -0.027041740
616   MPO -0.33275154
617   GCR 0.66710253
618 KRAS -1.54539394
619   TNF 0.55516440
620 TLR5 -1.50890306
621   FAS -0.63270525
622 FASL -0.36613984
623   P53 0.27965714
624 IKBKB -0.081503478
625 Case   CS6
626 TLR4 -0.52046012
627 TWIST1 0.89705746
628 TLR2 -1.72940907
629 TLR7 -1.48704529
630 NFKB1 -1.31309310
631 ELANE 0.712510907
632   MPO -1.49995383
633   GCR -1.13420338
634 KRAS -1.09756596
635   TNF -1.87275564
636 TLR5 -0.31439299
637   FAS 0.41996953
638 FASL 0.53631804
639   P53 -0.21611255
640 IKBKB -0.607721753
641 Case   CS7
642 TLR4 -0.13522719
643 TWIST1 0.56979981
644 TLR2 -0.28240124
645 TLR7 0.32297808
646 NFKB1 -1.72666511
647 ELANE -1.095487594
648   MPO -1.40809356
649   GCR -1.05866018
650 KRAS 0.67648840
651   TNF -1.65415954
652 TLR5 -1.06590585
653   FAS -0.80290371
654 FASL 1.02994108
655   P53 0.93078688
656 IKBKB 1.233548817
657 Case   CS8
658 TLR4 -0.46086511

659 TWIST1 0.11275207
660 TLR2 0.07167195
661 TLR7 -0.47729720
662 NFKB1 -0.49942986
663 ELANE -1.894723778
664   MPO 0.16832369
665   GCR -0.65574806
666 KRAS -0.14376994
667   TNF 0.41165935
668 TLR5 0.93723883
669   FAS 0.59149884
670 FASL -0.12560480
671   P53 -0.71474772
672 IKBKB 1.189507729
673 Case   CS9
674 TLR4 0.05137840
675 TWIST1 0.63661488
676 TLR2 -1.23140043
677 TLR7 -1.55721354
678 NFKB1 0.64903691
679 ELANE -0.796901744
680   MPO -0.40897787
681   GCR -0.81076882
682 KRAS 0.58927054
683   TNF 0.18523866
684 TLR5 -0.50168332
685   FAS -0.83418128
686 FASL 1.00426551
687   P53 -0.69722286
688 IKBKB 1.096094704
689 Case   CS10
690 TLR4 0.63720970
691 TWIST1 0.20227326
692 TLR2 -1.54234076
693 TLR7 -1.40871170
694 NFKB1 0.78359527
695 ELANE 0.005388105
696   MPO -0.67916926
697   GCR 0.24758493
698 KRAS 0.79749209
699   TNF -0.57404087
700 TLR5 0.28898534
701   FAS 1.06830448
702 FASL 1.03344012

703   P53   0.10573464
704 IKBKB 0.406900958
705   Case   CS11
706   TLR4   0.91570299
707 TWIST1   0.34058798
708   TLR2   0.79231152
709   TLR7   0.45723103
710 NFKB1 -0.21987084
711 ELANE 0.959635089
712   MPO   0.32735930
713   GCR   0.57685039
714   KRAS -1.20439277
715   TNF -0.60266507
716   TLR5   0.24930949
717   FAS -0.51940477
718   FASL   0.15869162
719   P53 -0.94189508
720 IKBKB -0.570522750
721   Case   CS12
722   TLR4 -0.98285515
723 TWIST1 -0.50508538
724   TLR2 -1.26040164
725   TLR7 -0.53851589
726 NFKB1   0.09973467
727 ELANE -1.623950480
728   MPO -0.87461525
729   GCR -0.47836682
730   KRAS   0.06831245
731   TNF   0.37901915
732   TLR5 -0.41087490
733   FAS   0.79090381
734   FASL -0.44944342
735   P53 -0.04419648
736 IKBKB -0.900111783
737   Case   NC1
738   TLR4 -0.87891891
739 TWIST1 -1.30618983
740   TLR2   0.37152646
741   TLR7 -1.17612846
742 NFKB1   0.23624269
743 ELANE -0.189718273
744   MPO -1.39407892
745   GCR   0.82905126
746   KRAS -1.90783721

747 TNF -1.02184732
748 TLR5 0.72832850
749 FAS 0.87156890
750 FASL 0.70256997
751 P53 1.15186593
752 IKBKB 0.213092722
753 Case   NC2
754 TLR4 -0.62565554
755 TWIST1 -1.90765719
756 TLR2 -1.71965026
757 TLR7 -0.45272755
758 NFKB1 -1.72189575
759 ELANE 0.551224969
760 MPO -0.69881195
761 GCR -1.60188065
762 KRAS -1.51884118
763 TNF 0.03957763
764 TLR5 -0.37183645
765 FAS 0.26909745
766 FASL -0.52980807
767 P53 1.25441649
768 IKBKB -0.006637377
769 Case   NC3
770 TLR4 -1.15376825
771 TWIST1 -1.57354893
772 TLR2 -0.91818135
773 TLR7 -1.85011531
774 NFKB1 0.19740610
775 ELANE -1.701890782
776 MPO -0.47747884
777 GCR -0.66000399
778 KRAS -1.95502358
779 TNF 0.38079496
780 TLR5 -0.68666972
781 FAS 0.25839576
782 FASL -0.85135032
783 P53 -0.35521648
784 IKBKB 1.275251247
785 Case   NC4
786 TLR4 -1.52555965
787 TWIST1 -1.12857708
788 TLR2 0.66251322
789 TLR7 -1.30263693
790 NFKB1 0.32108073

791 ELANE 0.899099471
792 MPO 0.88821532
793 GCR 0.77920350
794 KRAS 0.61235804
795 TNF -0.20768564
796 TLR5 -1.06608456
797 FAS -0.14359508
798 FASL 0.01540689
799 P53 -0.03875096
800 IKBKB 1.474491294
801 Case NC5
802 TLR4 0.56754905
803 TWIST1 -1.54958554
804 TLR2 -0.43901643
805 TLR7 -0.97015498
806 NFKB1 -1.55395991
807 ELANE 0.252097492
808 MPO -0.60330739
809 GCR -0.40120933
810 KRAS 0.09742885
811 TNF -0.20044751
812 TLR5 -1.56852562
813 FAS -0.73469211
814 FASL 0.58792331
815 P53 1.38061895
816 IKBKB 1.255136750
817 Case NC6
818 TLR4 -1.88813912
819 TWIST1 0.48116058
820 TLR2 -0.36080209
821 TLR7 0.19836664
822 NFKB1 -1.48759102
823 ELANE 0.456315948
824 MPO -1.59688306
825 GCR -0.09053130
826 KRAS 0.21881552
827 TNF -1.90593025
828 TLR5 -0.28653885
829 FAS -0.65574564
830 FASL -0.25996285
831 P53 0.97874239
832 IKBKB 1.372863318
833 Case NC7
834 TLR4 -0.35259296

835 TWIST1 -1.78195351
836 TLR2 0.80510938
837 TLR7 -1.85757360
838 NFKB1 0.24157416
839 ELANE -1.453589589
840   MPO -1.21589968
841   GCR 0.51937020
842 KRAS -1.38443083
843   TNF -0.64409009
844 TLR5 -0.15260244
845   FAS 1.36284886
846 FASL -0.32064879
847   P53 1.01125845
848 IKBKB -0.811012844
849 Case   NC8
850 TLR4 0.67058970
851 TWIST1 0.04054484
852 TLR2 -0.71318395
853 TLR7 0.04815973
854 NFKB1 -1.51101505
855 ELANE -1.572730650
856   MPO -0.35744430
857   GCR -1.57974926
858 KRAS -1.69696374
859   TNF 0.37842296
860 TLR5 -0.28756756
861   FAS 1.45513517
862 FASL 1.17727865
863   P53 -0.75886827
864 IKBKB 0.279848280