1	The: Differential anugenic requirements by diverse MR1-restricted 1 cens
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7	List of supplementary items:
8	Figure S1: Sequence verification of CRISPR/Cas9-mediated knockout of MR1 in 293T and
9	A375 cells
10	Figure S2: Effects on activation by 6-FP and Ac-6-FP
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15	Figure S1: Sequence verification of CRISPR/Cas9-mediated knockout of MR1 in 293T
16	and A375 cells. A. Alignment of nucleotide sequences of exon 3 of MR1 from position 15,885
17	to 15,944 from genomic DNA derived from 293T. ^{WT} , A375. ^{WT} , 293T.MR1 ^{KO} and
18	A375.MR1 ^{KO} cell lines. The coding sequence (CDS) is shown at the top and is derived from
19	NCBI reference ID NM_001531.3. Dots represent nucleotides identical to that of the CDS.
20	Dashes represent the absence of a nucleotide. The sequence of the sgRNA used during
21	CRISPR/Cas9-mediated knockout is highlighted in orange. Alignment was generated using
22	CLC Main Workbench 8 (Qiagen). B. Sanger Sequencing trace from genomic DNA amplified
23	from 293T.MR1 ^{KO} and A375.MR1 ^{KO} . The red dotted line indicated where a stretch of DNA
24	has been removed relative to the MR1 CDS. Traces were generated using 4Peaks software.
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Figure S2: Effects on activation by 6-FP and Ac-6-FP. A. Bar graphs showing activation of
TCR-expressing SKW-3 cells as measured by CD69, after co-culture with C1R.MR1^{HI} cells
B. Bar graphs showing the MFI of MR1 expression on A375.MR1^{HI} or A375.MR1^{K43A} cells
when pulsed with titrating concentrations of 6-FP or Ac-6-FP. Points on all plots are replicate
wells, and data is representative of n=3 independent experiments.

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Figure S3: MR1 tetramer staining of MR1-restricted TCRs. Flow cytometric contour plots
showing MR1-5-OP-RU and MR1-6-FP tetramer staining on 293T cells transiently transfected
to express diverse MR1-restricted TCRs. Experiment is representative of 2 independent
experiments.

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Figure S4: Folate-free media has no effect on activation of MR1-restricted TCRs. Bar graphs showing activation of TCR-expressing SKW-3 cells as measured by CD69, after coculture with a panel of antigen-presenting cell lines that had been cultured for 48 hours in normal RF10 media, or RF10 media that used folate-free RPMI-1640 as a base. Points on plots are replicate wells, and data is representative of n=2 independent experiments.

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Figure S5: TCR tetramers fail to stain MR1-overexpressing cell lines. A. ELISA showing binding of anti- $\alpha\beta$ TCR, anti-V α 7.2 or anti- $\gamma\delta$ TCR to soluble TCR proteins. **B-D.** Flow cytometric contour plots showing TCR tetramer staining on **B.** C1R cell lines, **C.** A375 cell lines, or **D.** A375.MR1 cell lines pulsed for 4 hours with 10 µM 5-OP-RU, 500 µM Ac-6-FP or 500 µM 6-FP. Plots are representative of n=2 independent experiments. Numbers in top right of plots indicate MFI of tetramer staining.

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Point of KO



6-FP Ac-6-FP

6-FP Ac-6-FP





