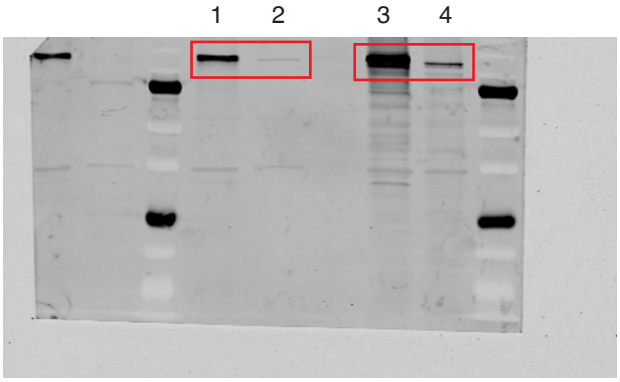
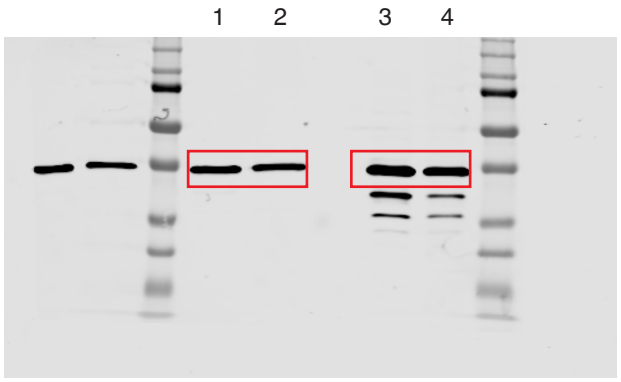
a

Blot 1
RB1

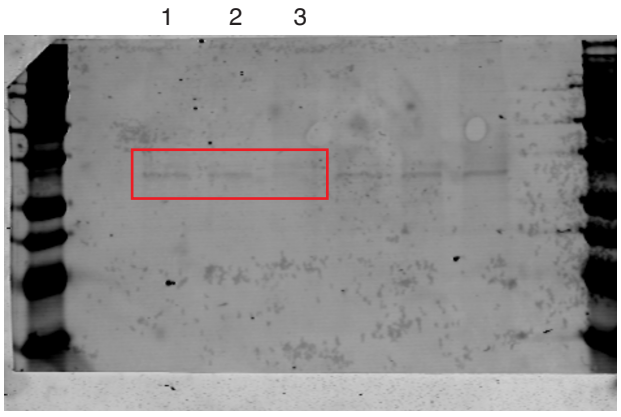


GAPDH

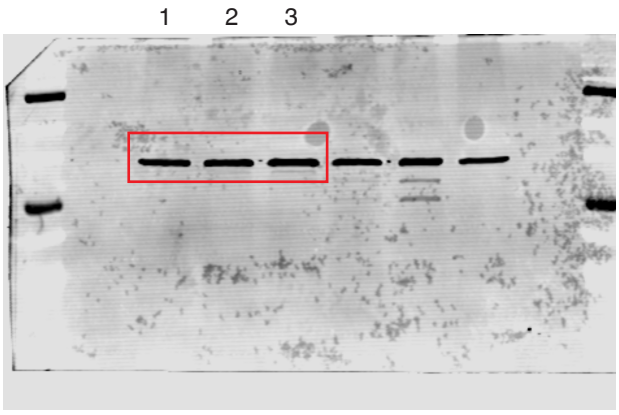


b

Blot 2
TREX1

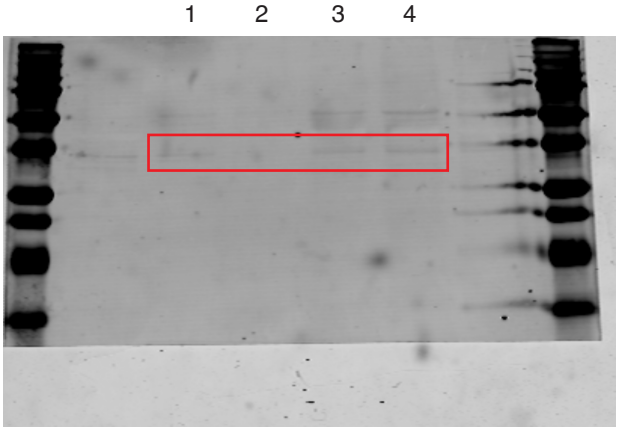


GAPDH

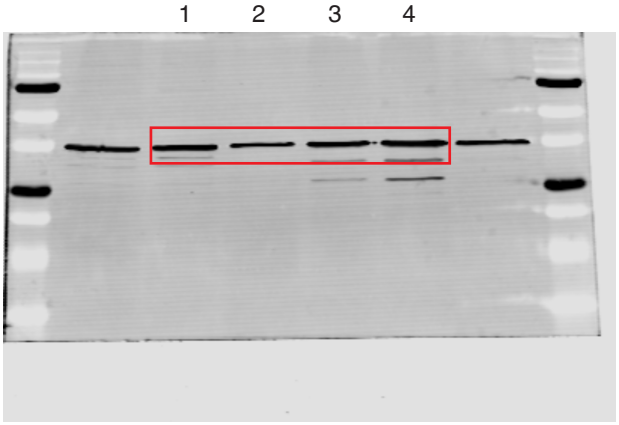


b

Blot 3
FOXA1

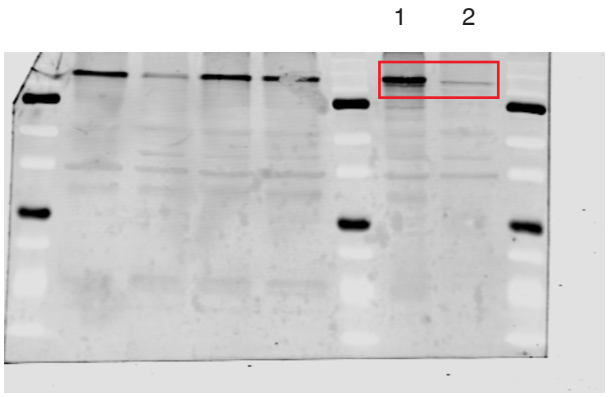


GAPDH

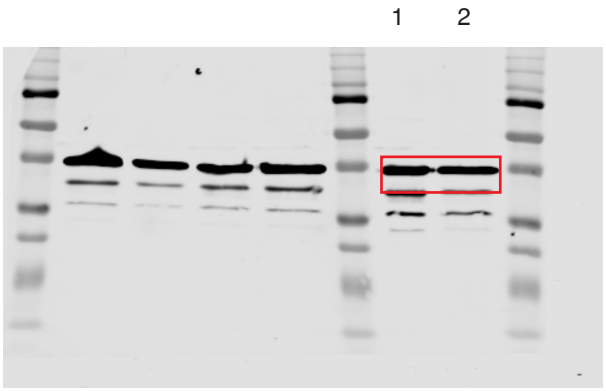


d

Blot 4
FOXA1

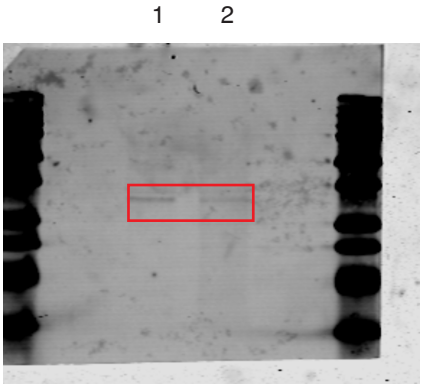


GAPDH

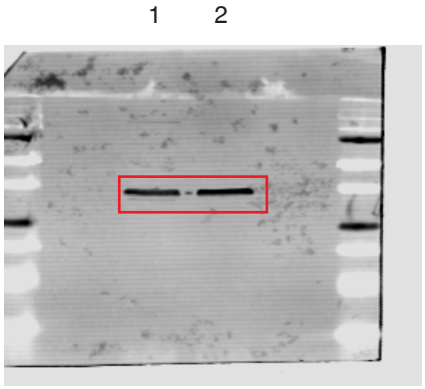


e

Blot 5
TREX1

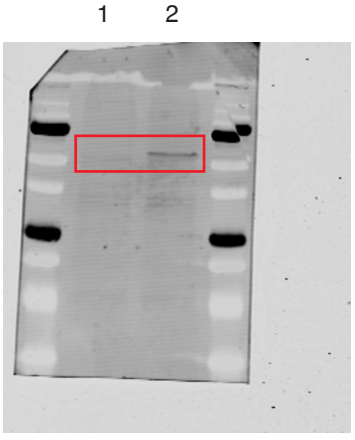


GAPDH

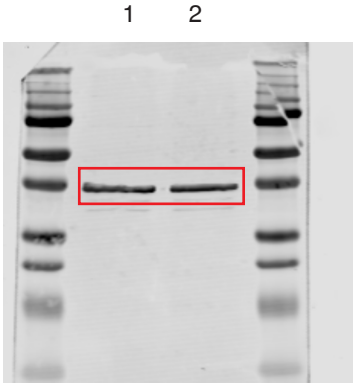


f

Blot 6
FOXA1

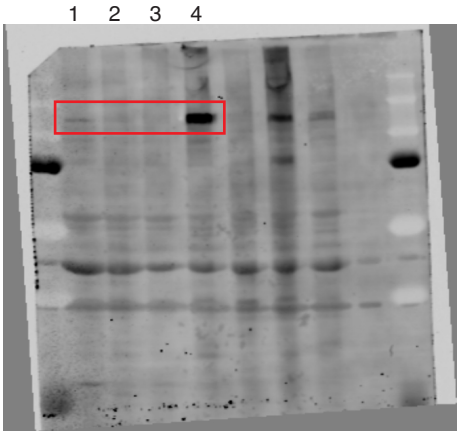


GAPDH



g

Blot 7
RB1



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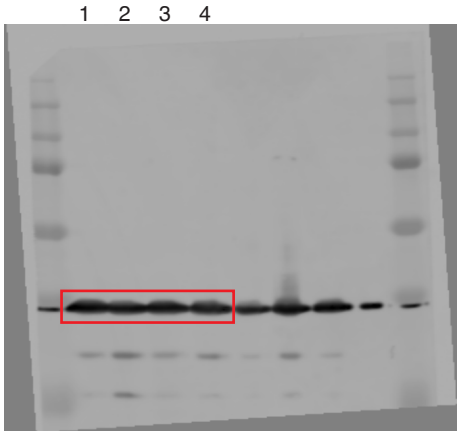
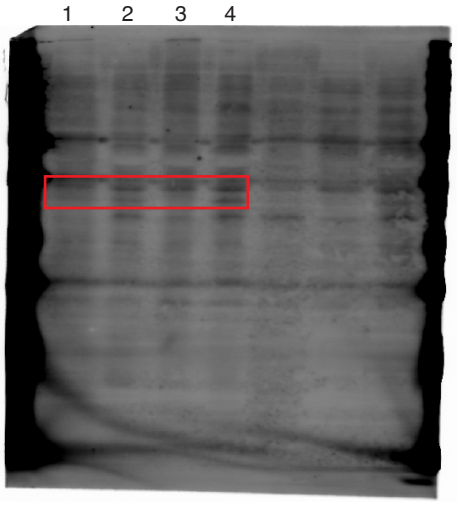


h

Blot 8
TREX1



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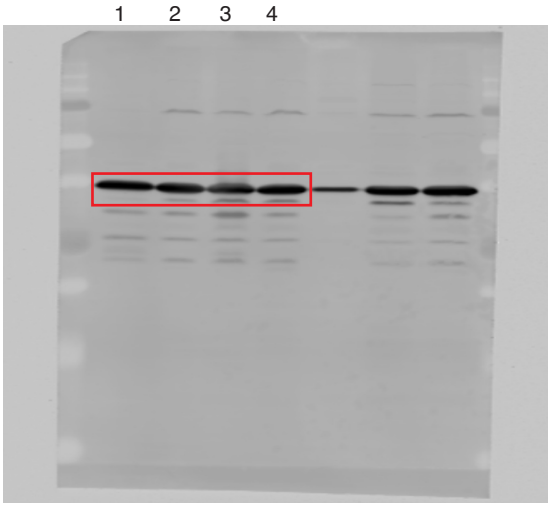
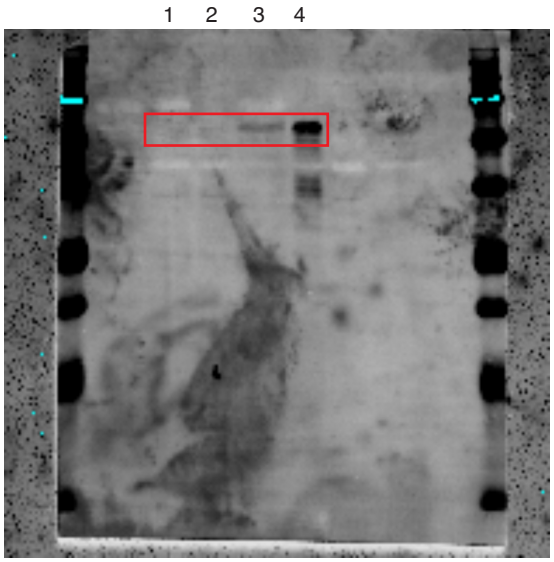
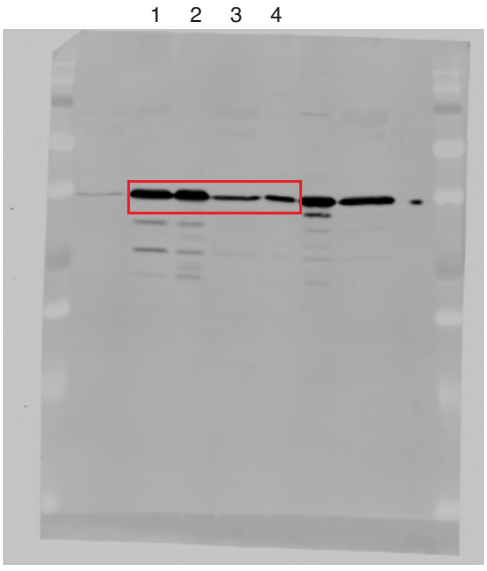


i

Blot 9
FOXA1

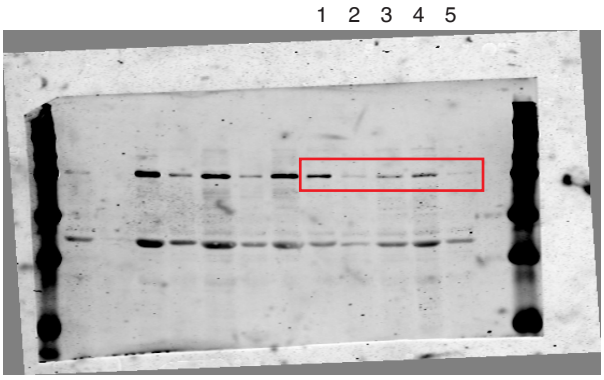


GAPDH

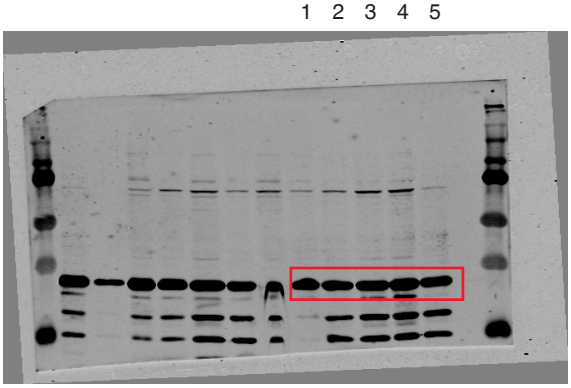


j

Blot 10
STAT2

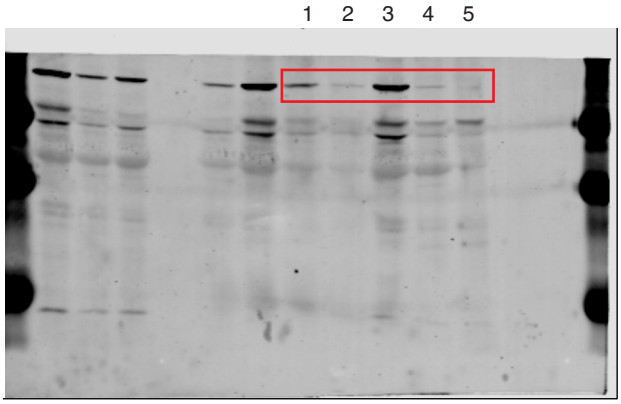


GAPDH

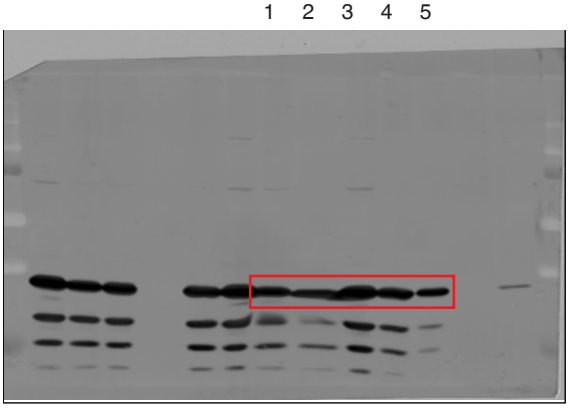


k

Blot 11
IRF7

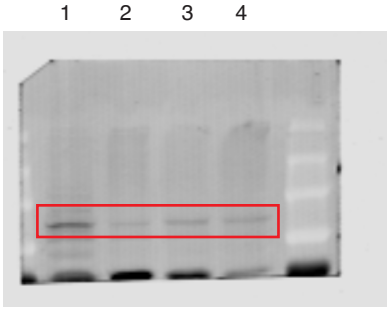


GAPDH

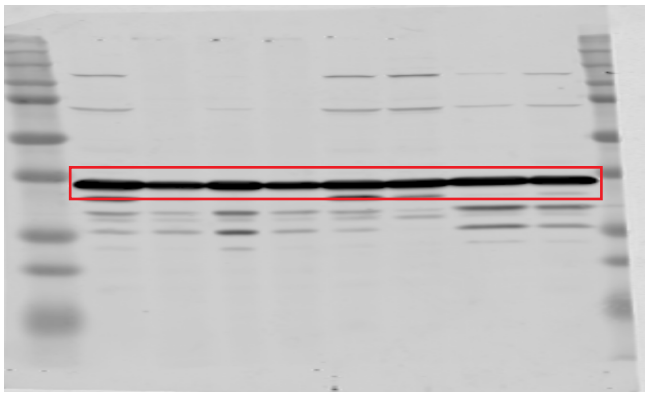


l

Blot 12
STAT2

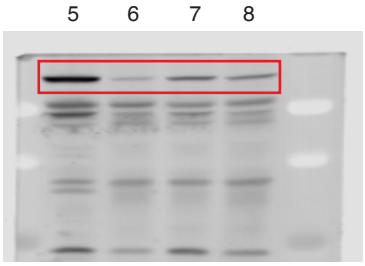


GAPDH

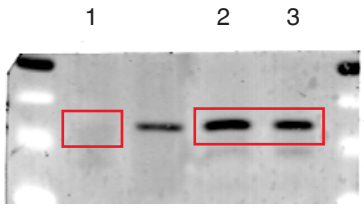


m

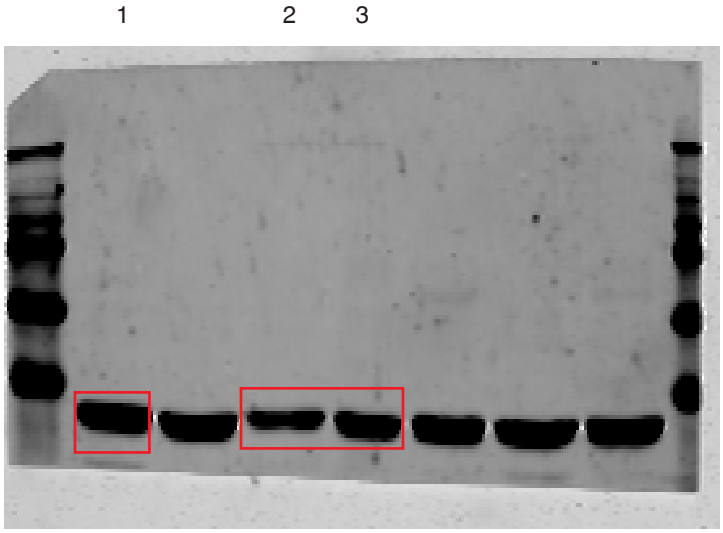
Blot 13
IRF7



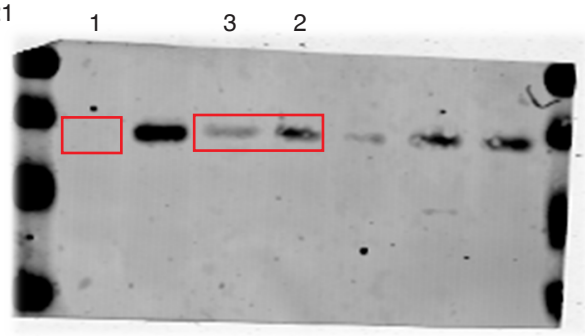
n Blot 14
p16



GAPDH



o Blot 15
p21



a, Blot 1 - RB1: 1. EP; 2. SEN (L); 3. OE – SEN; 4. SEN (E)

b, Blot 2 - TREX1: 1. EP; 2. SEN (E); 3. SEN (L)

c, Blot 3 - FOXA1: 1. EP; 2. Arrest; 3. SEN (E); 4. SEN (L)

d, Blot 4 - RB1: 1. EP; 2. 3X

e, Blot 5 - TREX1: 1. EP; 2. 3X

f, Blot 6 - FOXA1: 1. EP; 2. 3X

g, Blot 7 - RB1: 1. SEN (L); 2. shRB1(b); 3. shRB1(a); 4. OE-RB1

h, Blot 8 - TREX1: 1. SEN (L); 2. shTREX1 (b); 3. shTREX1 (a); 4. OE-TREX1

I, Blot 9 - FOXA1: 1. shFOXA1 (a); 2. shFOXA1 (b); 3. SEN (L); 4. OE-FOXA1

j, Blot 10 - STAT2: 1. 3X; 2. shL1; 3. NRTI; 4. ΔIFNAR; 5. EP

k, Blot 11 - IRF7: 1. EP; 2. shL1; 3. 3X; 4. shL1; 5. ΔIFNAR

l, Blot 12 - STAT2: 1. SEN (L); 2. ΔIFNAR; 3. shL1; 4. NRTI

m, Blot 13 - IRF7: 5. SEN (L); 6. ΔIFNAR; 7. shL1; 8. NRTI

n, Blot 14 – p16 (CDKN2A): 1. EP; 2. SEN (E); 3. SEN (L)

o, Blot 15 – p21 (CDKN1A): 1. EP; 2. SEN (E); 3. SEN (L)