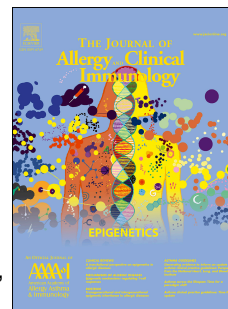


Accepted Manuscript

EROS mutations: decreased NADPH oxidase function and chronic granulomatous disease

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1 **Full Title: : EROS mutations: decreased NADPH oxidase function and chronic**
2 **granulomatous disease**

3

4 **Running Title: *EROS* mutations cause chronic granulomatous disease**

5

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26

27 **Conflict of interest statement**

28 The authors have no conflicts of interest to declare.

29

30 **Capsule Summary**

31 We demonstrate for the first time that EROS (CYBC1/C17ORF62) regulates abundance of
32 the gp91phox-p22phox heterodimer of the phagocyte NADPH oxidase in human cells and
33 that EROS mutations are a novel cause of chronic granulomatous disease.

34

35

36 **Key Words**

37 EROS,

38 *C17ORF62*39 *CYBC1*

40 Chronic granulomatous disease,

41 Nox2

42 *gp91phox*

43

44 **Abbreviations**

45 CGD – Chronic Granulomatous Disease

46 EROS – Essential for Reactive Oxygen Species

47 HLH – haemophagocytic lymphohistiocytosis

48 NADPH - Nicotinamide adenine dinucleotide phosphate

49 NBT Nitro blue tetrazolium chloride

50 DHR – Dihydrorhodamine

51 PBMC - Peripheral Blood Mononuclear Cell

52

53

54 **Short Summary (for the Editor only)**

55 The phagocyte respiratory burst is mediated by the phagocyte NADPH oxidase, a multi-
56 protein subunit complex that facilitates production of reactive oxygen species and which is
57 essential for host defence. Monogenic deficiency of individual subunits leads to chronic
58 granulomatous disease (CGD), which is characterized by an inability to make reactive
59 oxygen species, leading to severe opportunistic infections and auto-inflammation. However,
60 not all cases of CGD are due to mutations in previously identified subunits. We recently
61 showed that Eros, a novel and highly conserved ER-resident transmembrane protein, is
62 essential for the phagocyte respiratory burst in mice because it is required for expression of
63 gp91*phox*-p22*phox* heterodimer, which are the membrane bound components of the
64 phagocyte NADPH oxidase. Eros has a human orthologue, *CYBC1/EROS*. We now show that
65 the function of *CYBC1/EROS* is conserved in human cells and describe a case of CGD
66 secondary to a homozygous *CYBC1/EROS* mutation that abolishes EROS protein expression.
67 This work demonstrates the fundamental importance of *CYBC1/EROS* in human immunity
68 and describes a novel cause of CGD.

69

70 **To the editor:**

71 The multi-subunit phagocyte NADPH oxidase generates reactive oxygen species and is
72 crucial for host defence (1). Deficiencies in individual subunits (gp91*phox*, p22*phox*,
73 p47*phox*, p67*phox* and p40*phox*) cause chronic granulomatous disease (CGD) but some
74 patients with CGD do not have mutations in these genes (2). We recently found that Eros (3),
75 a hitherto undescribed protein, is essential for the generation of reactive oxygen species
76 because it is necessary for protein (but not mRNA) expression of the gp91*phox*-p22*phox*
77 heterodimer, which is almost absent in *Eros*-deficient mice. *Eros*^{-/-} animals succumb quickly
78 following infection with *Salmonella* Typhimurium or *Listeria Monocytogenes*. *Eros* is highly

79 conserved and has a human orthologue *CYBC1* (alias *C17ORF62*), hereafter referred to as
80 *CYBC1* gene and EROS protein). We asked whether the gene fulfilled the same function in
81 humans. We performed CRISPR-mediated deletion of *CYBC1/EROS* in PLB-985 cells, (**Fig.**
82 **S1A**) and identified two clones with 8bp and 1bp deletions respectively (**Fig. S1B and C**).
83 Neither clone expressed EROS protein (**Fig. 1A**) or detectable *gp91phox* (**Fig. 1B**). *p22phox*
84 expression was also much lower in both *EROS*-deficient clones than in control cells (**Fig.**
85 **1C**). We verified the lack of surface *gp91phox* expression by flow cytometry (**Fig. 1D**). Both
86 *CYBC1/EROS*-deficient clones had a severely impaired respiratory burst (**Fig. 1E**). In
87 addition, *CYBC1/EROS*-deficient clones differentiated towards a neutrophil phenotype also
88 demonstrated an impaired respiratory burst (**data not shown**). As expected, re-introduction
89 of *CYBC1/EROS* using a lentiviral vector restored *gp91phox* expression to *CYBC1/EROS*-
90 deficient clones (**Figure 1F**) and oxidase activity as measured by Nitro blue tetrazolium
91 chloride (NBT) test (**Figure 1G**) and DIOGENES assay (data not shown).

92
93 We then identified a patient with a homozygous *CYBC1/EROS* mutation in a resource paper
94 that details a thousand Saudi Arabian families with genetic disease(4). He presented with
95 fever, splenomegaly, lymphadenopathy and short stature, but no immuno-phenotyping was
96 detailed at that time. His full clinical history is as follows. He is a Saudi Arabian boy, born in
97 2007, the son of parents in a consanguineous marriage. He has three healthy older sisters. At
98 2 months of age, he developed a localized abscess following BCG vaccination. He was then
99 relatively well until 8 years of age but was noted to be of short stature and experienced
100 recurrent pulmonary infections and tonsillitis/pharyngitis despite tonsillectomy.

101

102 In August 2015, he became unwell with a febrile illness and an abnormal dihydrorhodamine
103 (DHR) test was noted (**Fig. 2A,B**). He has a severely impaired DHR in response to both PMA

104 and zymosan. He was also profoundly lymphopenic. He subsequently developed an acute
105 episode of hemolytic anemia in November 2015 and again in January 2017. During 2015, his
106 fevers, infection, lymphopenia and elevated inflammatory markers met criteria for a
107 diagnosis of haemophagocytic lymphohistiocytosis (HLH). He had no mutations previously
108 implicated in HLH pathogenesis. He was treated with cyclosporine and steroids and was
109 transferred to Boston Children's Hospital (BCH) in December 2016 for consideration of bone
110 marrow transplantation. The DHR test was repeated and rhodamine fluorescence was again
111 virtually absent. Further assessment demonstrated granulomatous inflammation in his lungs,
112 and discrete granulomata in his bone marrow, with no evidence of infection. Following his
113 open lung biopsy at BCH, he developed hemolytic anemia and required Intensive Care. He
114 recovered with steroids, however on weaning of this therapy he developed recurrent pleural
115 effusions. Due to his autoimmunity, features of lymphopenia with granulomata, and pleural
116 effusions/hemolytic anemia he was started on sirolimus and he remains in this therapy. His
117 steroids have been weaned to 4mg daily. While reasonably well clinically, he developed
118 diminished anti-pneumococcal antibody responses, worsening lymphopenia, and declining
119 immunoglobulin levels. He therefore underwent a myeloablative bone marrow transplant and
120 has recovered well.

121

122 Whole exome sequencing demonstrated that the patient had a homozygous (c.127 A to G,
123 NM_001033046) mutation in *CYBC1/EROS*. His sisters were all heterozygous for this
124 mutation, confirmed by Sanger sequencing (**Fig. 2C**). Based on analysis of other family
125 members and the likely important role of *CYBC1/EROS* in immunity, this mutation was
126 identified as the most likely cause of the patient's disease. The mutation was not present in
127 10,000 whole genomes from the United Kingdom National Institute for Health Research
128 BioResource - Rare Disease cohort (which includes 1000 patients with primary

129 immunodeficiency), nor in gnomAD. The variant was also absent in 3,300 ethnically
130 matched exomes. It is, therefore, not seen in seen across 110,579 individuals with sequence
131 data coverage across this position. There were no deleterious mutations in known NADPH
132 oxidase subunits.

133

134 Splice site prediction algorithms including Mutation Tasting (www.mutationtaster.org) and
135 Human Splicing Finder (<http://www.umd.be/HSF3/>) predict that the variant both disrupts an
136 exonic splice enhancer (ESE) and creates an exonic splice silencer (ESS) which is likely to
137 lead to a retained intron. This intron has 4 in-frame stop codons. It is therefore likely that
138 translation would cease downstream of exon 4. Even if splicing is not disrupted, PolyPhen
139 (<http://genetics.bwh.harvard.edu/pph2/>) predicts that the D43N mutation that would occur
140 in the translated protein is also damaging.

141

142 We therefore performed western blot analysis on anti-CD3-CD28-CD2 expanded T cells
143 from the patient, his sister and a healthy control, as well as primary T cells (either pre or post
144 polyclonal stimulation) and peripheral blood mononuclear cells from healthy volunteers. The
145 CRISPR targeted clones described above were used as positive and negative controls
146 respectively. The patient had undetectable levels of EROS protein compared with cells from
147 the healthy control or the primary T cells/PBMC, while his heterozygous sister had
148 intermediate levels (**Fig. 2D**).

149

150 This work demonstrates that the function of the novel transmembrane protein, Eros, is
151 conserved in humans. It also represents the first description of an immunodeficiency
152 syndrome secondary to mutations in *CYBC1/EROS*. The severity of the disease seen in this
153 patient underlines the importance of human EROS is for normal immunity. The patient has a

154 clinical history that is compatible with a diagnosis of autosomal recessive CGD, in that he
155 had both infectious and auto-inflammatory manifestations, together with histo-pathological
156 evidence of granuloma formation in the context of an impaired DHR response. While
157 recurrent infections, BCG-it is (2, 5), granulomatous inflammation and HLH (6) are all
158 recognised sequelae of CGD, this patient some unusual features such as autoimmune
159 haemolytic anaemia. This is uncommon in CGD and may represent an effect of EROS-
160 deficiency that is independent of its effects on the NAPDH oxidase.

161

162 The high sequence similarity between mouse and human EROS and the loss of *gp91phox*
163 expression and the phagocyte respiratory burst that accompanies its absence suggests that
164 EROS plays an almost identical role in human and murine immunity. In summary, we have
165 shown that the function of EROS is fully conserved between human and mouse, and that
166 homozygous mutations in EROS underlie a novel sixth cause of chronic granulomatous
167 disease.

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254

255 **Author Contributions**

256 DCT designed and performed experiments and wrote the manuscript. LC analysed patient
257 samples. **A Schejtman** designed guide RNAs, performed CRISPR-mediated deletion of
258 EROS in PLB-985 cells and NBT assays. EC prepared plasmids for transfection experiments
259 and cultured cells. LD performed western blots and assisted with culture of the PLB-985
260 cells. AB assisted with western blots and ROS assays. JL advised on and performed
261 electroporation of PLB-985 cells. SC and A.Speak advised on cloning strategies and oversaw
262 plasmid preparation. AT and GS oversaw CRISPR-mediated deletion experiments and
263 advised on other experiments. FSA co-ordinated the sequencing of the patient and provided
264 advice on experiments. HA and HAM treated and diagnosed the patient in Saudi Arabia. TC
265 oversaw the clinical care of the patient in Boston and contributed to writing the manuscript.
266 KGCS oversaw experiments in the project and wrote the manuscript.

267

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284

285 **Figure 1: EROS function is conserved in humans.** (A-C) Western blot of (A) EROS (B)
 286 gp91*phox* (C) p22*phox*, (D) surface gp91*phox* expression and (E) phagocyte respiratory
 287 burst in CRISPR targeted PLB-985 clones (F) surface expression of gp91*phox* in CRISPR
 288 targeted cells +/- overexpression of EROS-GFP. (G) NBT reduction following reconstitution
 289 with EROS-GFP lentiviral vector. Representative of 3 independent experiments.

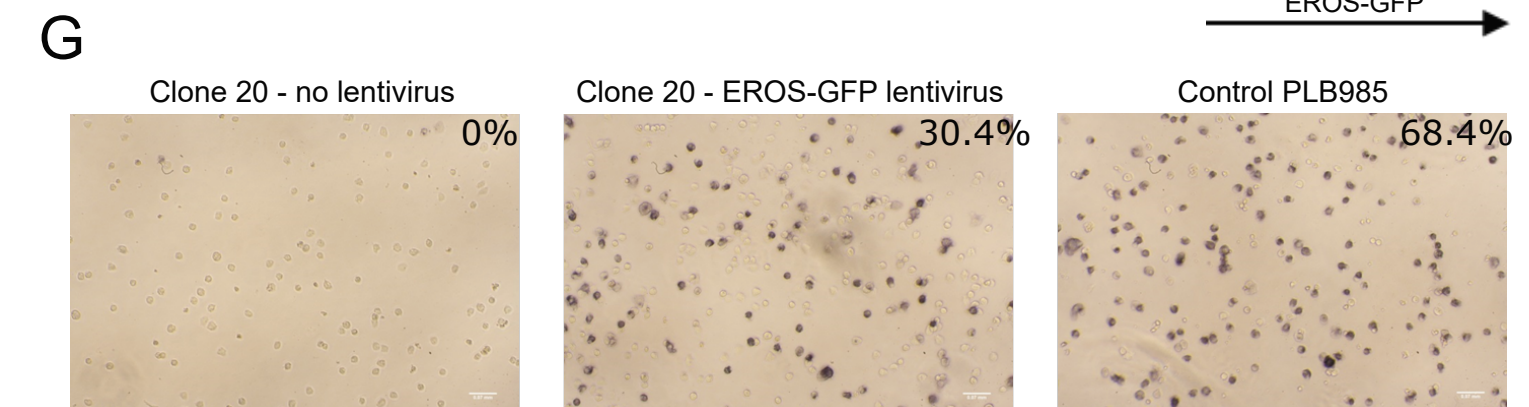
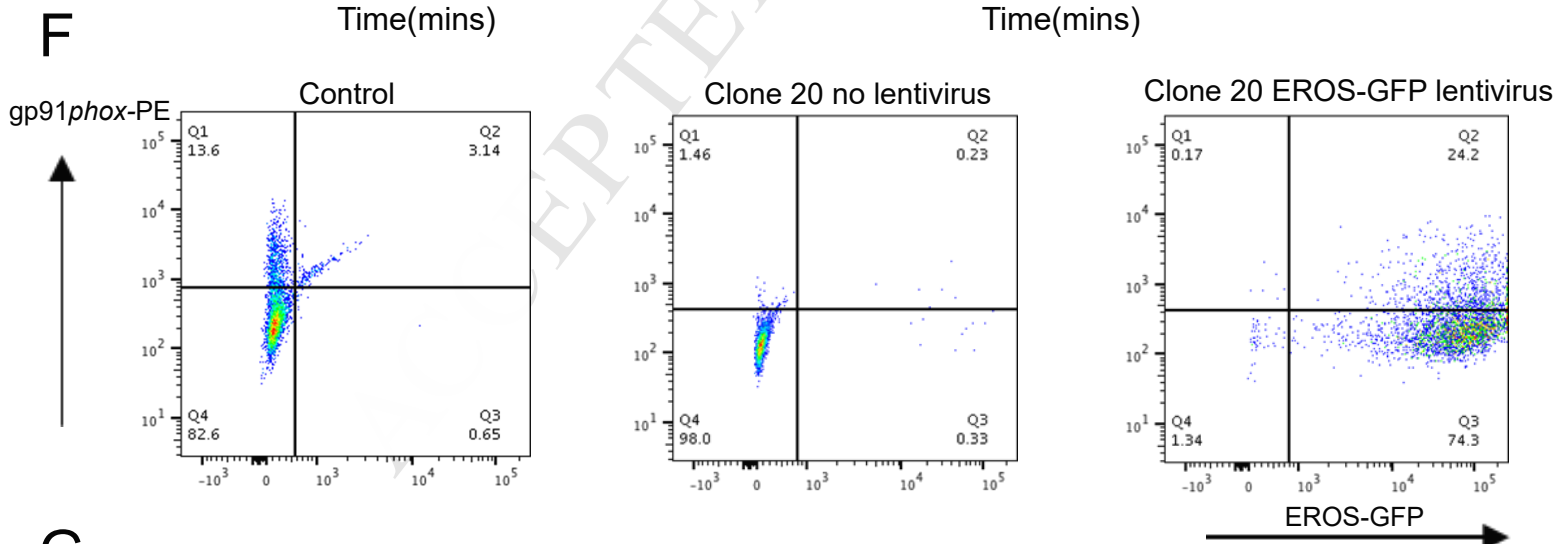
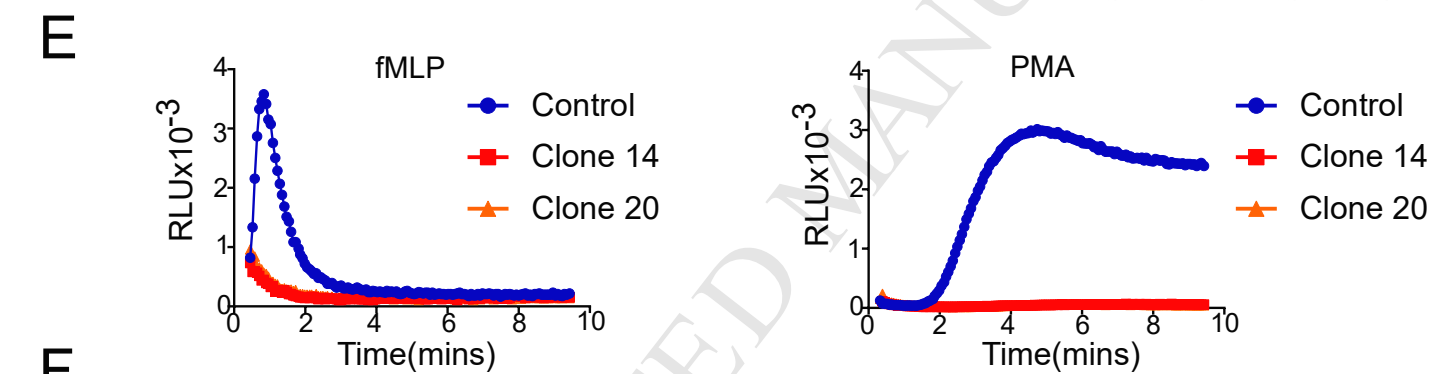
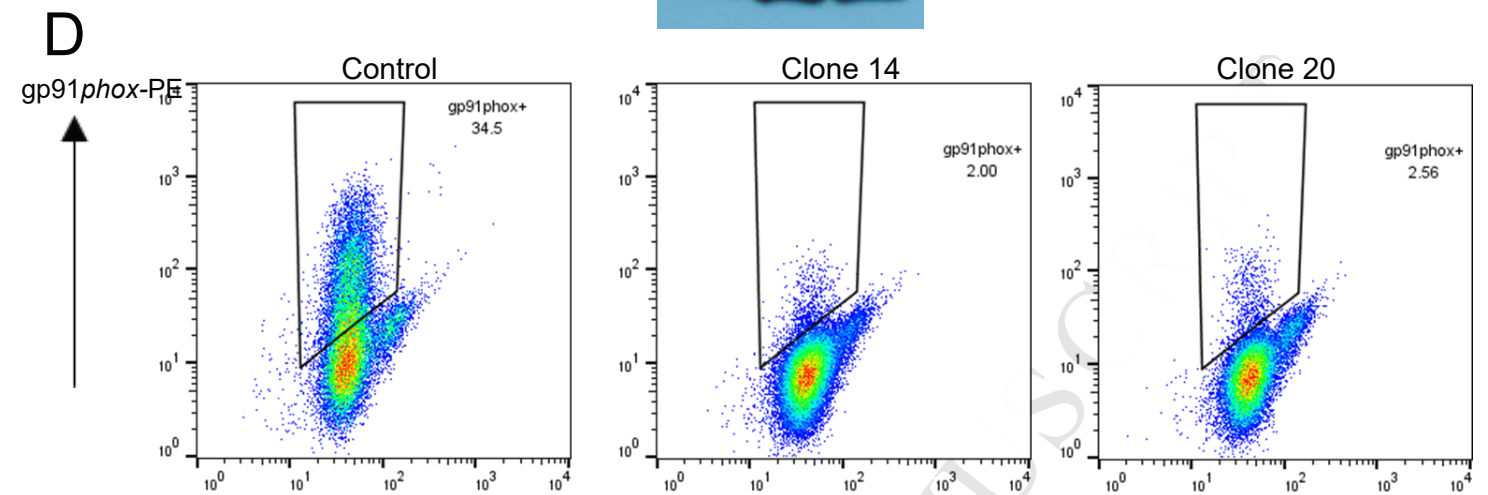
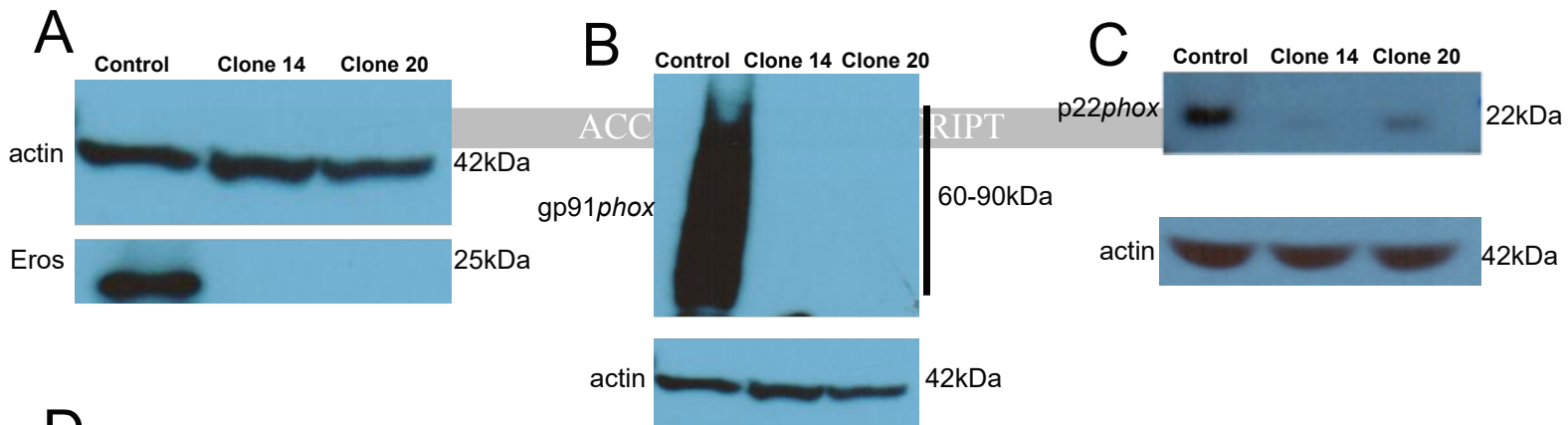
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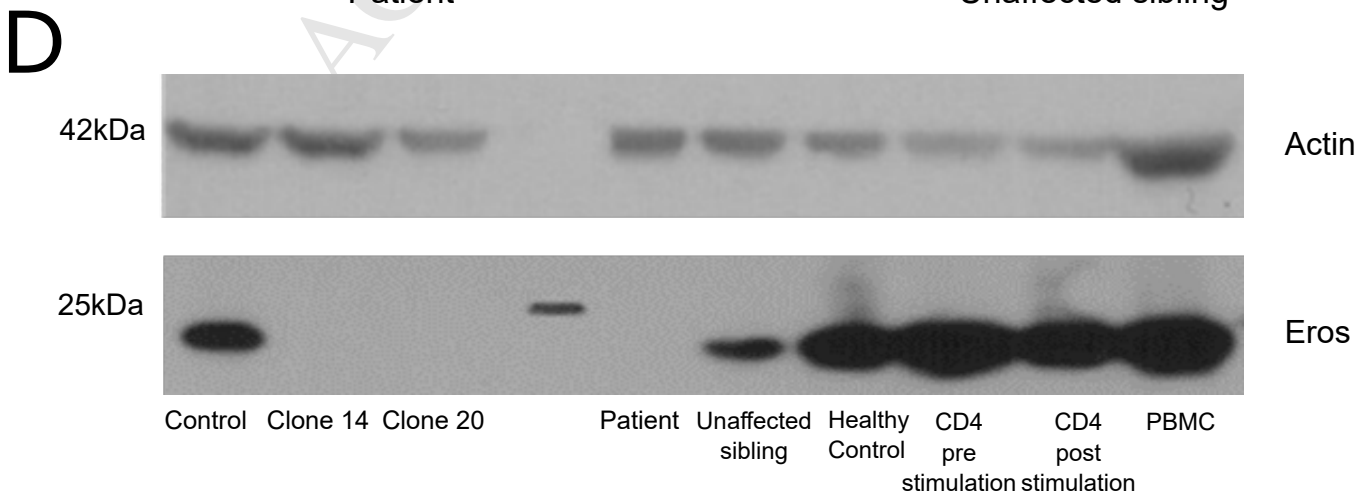
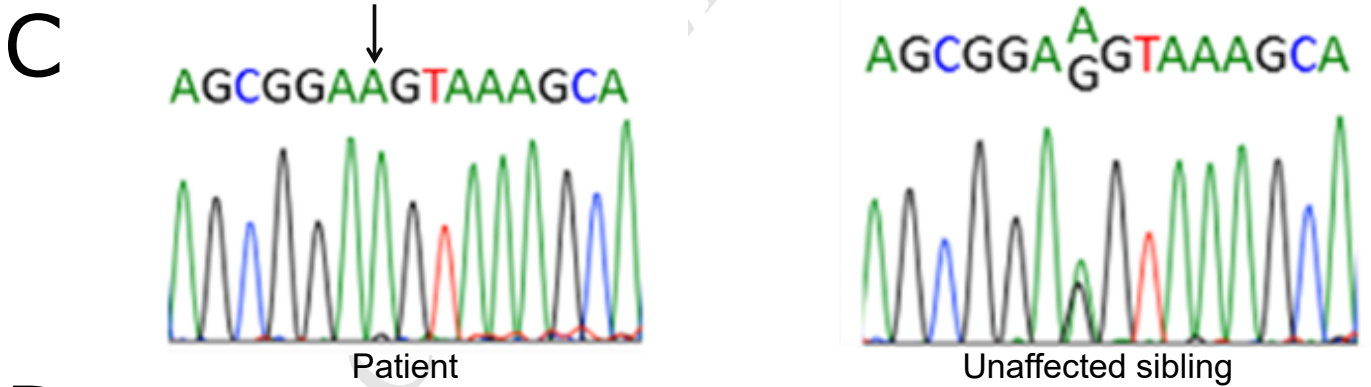
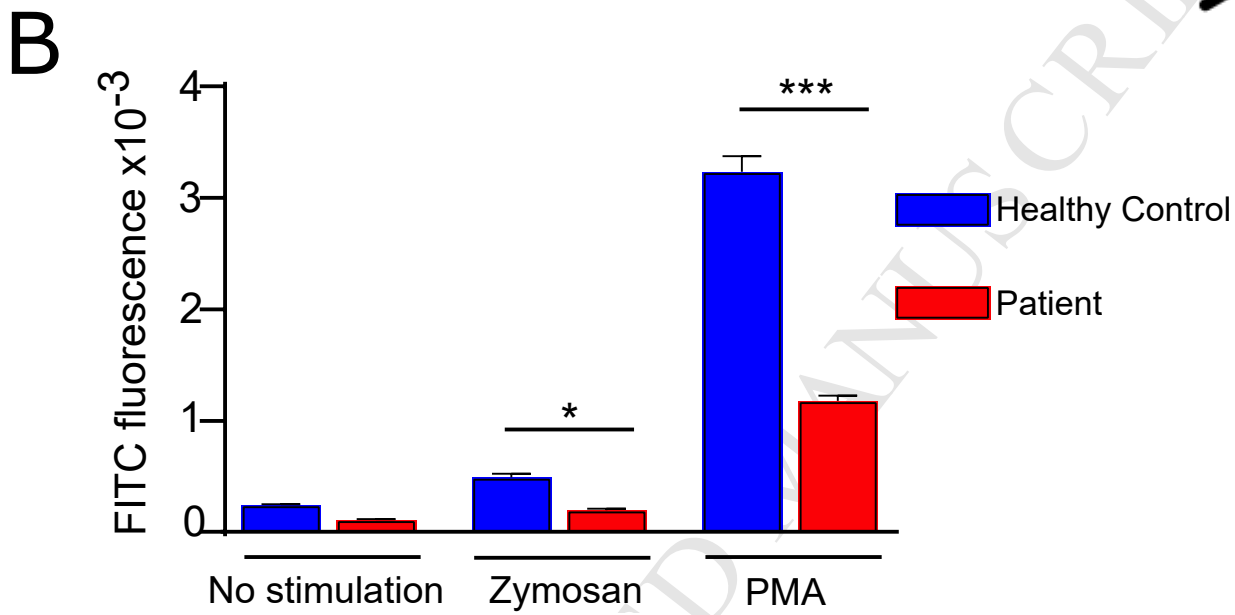
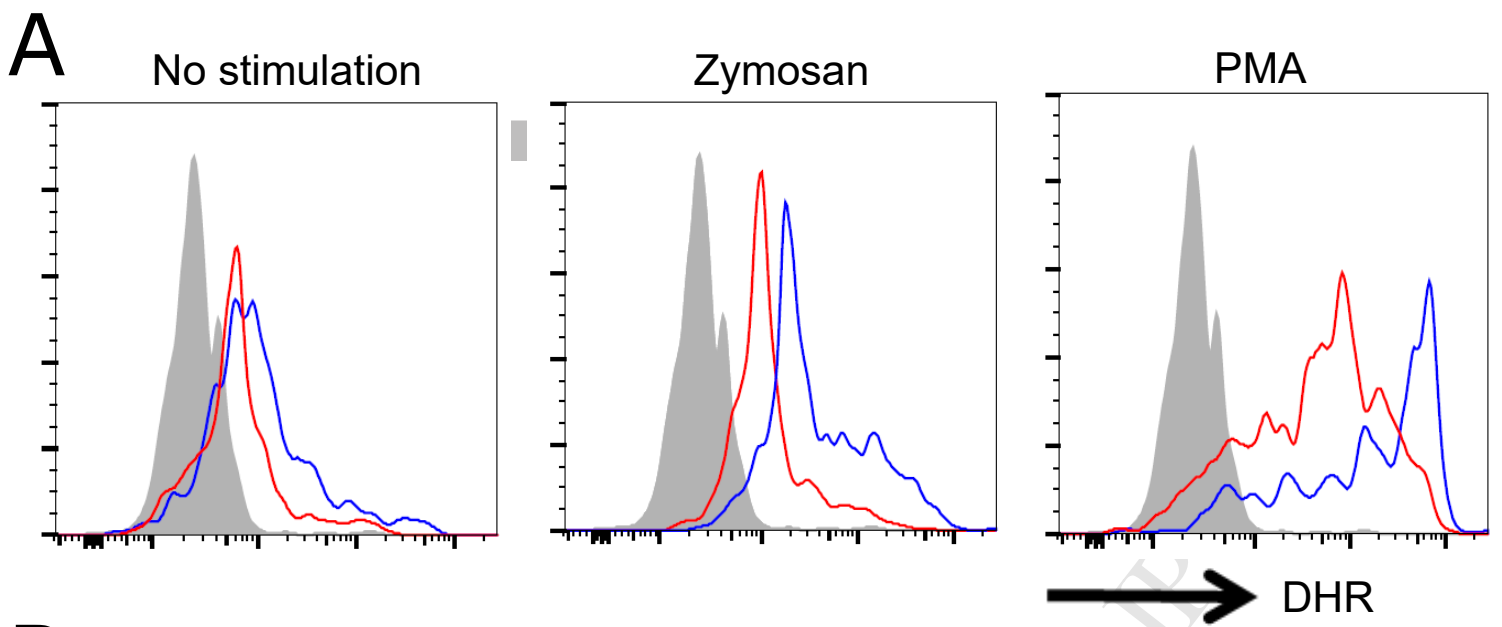
291 **Figure 2: Homozygous EROS mutation has deleterious effects.** (A, B): respiratory burst in
 292 patient and healthy control neutrophils. (C) homozygous c.127 A to G mutation (D) EROS
 293 protein expression in CD3/CD2/CD28 expanded PBMC from patient, sister and a healthy
 294 control, control CD4+ T cells pre and post CD3/CD2/CD28 or PBMC from a further healthy
 295 control. Representative of 2-3 independent experiments.

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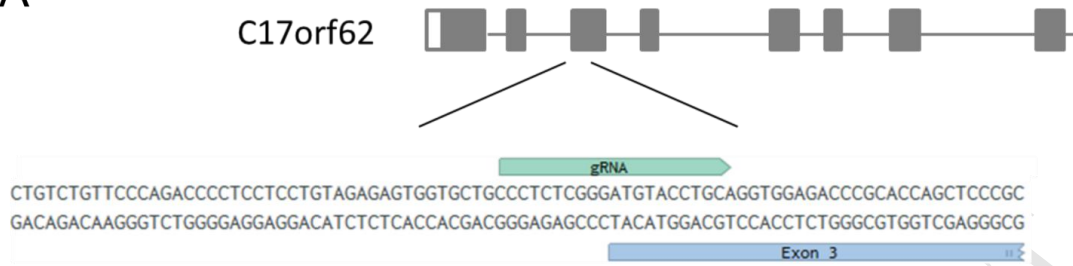


Legend for supplementary figure

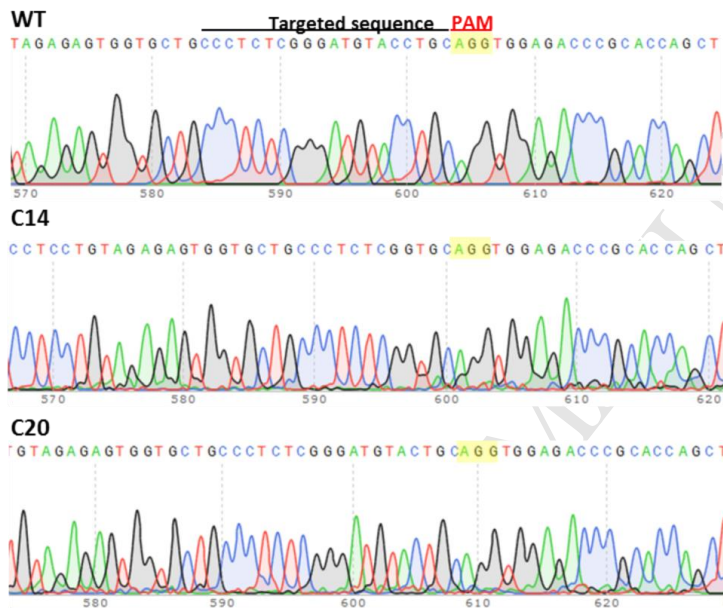
Figure S1: Targeting of EROS for deletion by CRISPR-Cas9 (A) Schematic representation of the human *CYBC1* genomic locus and of guide RNA targeting exon 3 (B, C) Sequencing analysis from two clones derived from PLB985 cells that had been electroporated with Cas9:gRNA ribonucleoprotein complex, showing deletion of 8bp in clone 14 and deletion of 1 bp in clone 20.

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A



B



C

8bp deletion

WT sequence CTGTTCCCAGACCCCTCCTCCTGTAGAGAGTGGTGTGCCCTCTCGGGATGTACCTGCAG

C14 sequence CTGTTCCCAGACCCCTCCTCCTGTAGAGAGTGGTGTGCCCTCTCGG-----TGAG

WT sequence GTGGAGACCCGCACCAGCTCCCGCTCCATCTGAAGAGGGTCCAGGCATCCGGTCTTG

C14 sequence GTGGAGACCCGCACCAGCTCCCGCTCCATCTGAAGAGGGTCCAGGCATCCGGTCTTG

1bp deletion

WT sequence CTGTCTGTTCCCAGACCCCTCCTCCTGTAGAGAGTGGTGTGCCCTCTCGGGATACCT

C20 sequence CTGTCTGTTCCCAGACCCCTCCTCCTGTAGAGAGTGGTGTGCCCTCTCGGGATCTA-C

WT sequence GCAGGTGGAGACCCGCACCAGCTCCCGCTCCATCTGAAGAGGGTCCAGGCATCCGGTC

C20 sequence GCAGGTGGAGACCCGCACCAGCTCCCGCTCCATCTGAAGAGGGTCCAGGCATCCGGTC