

Oxidative stress in the haematopoietic niche regulates the cellular immune response in *Drosophila*

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Oxidative stress induced by high levels of reactive oxygen species (ROS) is associated with the development of different pathological conditions, including cancers and autoimmune diseases. We analysed whether oxidatively challenged tissue can have systemic effects on the development of cellular immune responses using *Drosophila* as a model system. Indeed, the haematopoietic niche that normally maintains blood progenitors can sense oxidative stress and regulate the cellular immune response. Pathogen infection induces ROS in the niche cells, resulting in the secretion of an epidermal growth factor-like cytokine signal that leads to the differentiation of specialized cells involved in innate immune responses.

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INTRODUCTION

Studies during the past decade have shown an important role of oxidative stress sensing in the development and function of different tissues in metazoans (Finkel & Holbrook, 2000; Ghaffari, 2008; Owusu-Ansah *et al*, 2008; Johannsen & Ravussin, 2009). While most investigations were primarily centred on the cell autonomous effects of oxidative stress, the question of whether the stressed cells can generate signals that mediate cell non-autonomous effects on neighbouring or distant tissues has not been properly addressed. This question is of particular importance for the regulation of innate and adaptive immune responses and has possible involvement in the aetiology of immunological and haematopoietic diseases. In this paper we have investigated cell-non-autonomous effects caused by oxidative stress that alters the differentiation status of the stress-sensing myeloid-like *Drosophila* blood system.

A haematopoietic organ of *Drosophila* known as the lymph gland consists of three developmentally distinct cellular zones (Jung *et al*, 2005). Haematopoietic progenitor cells reside in a medullary zone (MZ) of the organ. During development, these progenitors differentiate to give rise to mature blood cells that populate the cortical zone. A cluster of cells located in the posterior end of the gland represents its third compartment, called the posterior signalling centre (PSC) that primarily functions as a haematopoietic niche or microenvironment, maintaining the stem-like progenitor cells within the MZ (Krzemien *et al*, 2007; Mandal *et al*, 2007). The cellular component of the *Drosophila* innate immune system is constituted of three types of effector blood cell: plasmatocytes, crystal cells and lamellocytes. Plasmatocytes are responsible for phagocytosis/scavenging and crystal cells are responsible for blood clotting; they are maintained in relatively constant numbers during development (Lanot *et al*, 2001). In contrast, lamellocytes are barely detectable under normal conditions but are rapidly generated on infestation by parasitic intruders such as wasp eggs, and these cells are responsible for the inactivation and removal of the invaders (Rizki & Rizki, 1992; Sorrentino *et al*, 2002; Markus *et al*, 2005). Lamellocyte differentiation induced by immune challenge fails to occur in the absence of *collier* expression in the PSC (Crozatier *et al*, 2004), suggesting the possibility that a signal from an intruder event mediates changes in the PSC, which in turn sends a signal to initiate differentiation of lamellocytes. In this study we have addressed the mechanism by which this systemic response is generated.

RESULTS AND DISCUSSION

High ROS in the PSC induces cellular immune response

Abnormal metabolism is often associated with oxidative stress that results in increased production of ROS by mitochondria (Finkel & Holbrook, 2000; Owusu-Ansah *et al*, 2008; Starkov, 2008). Different concentrations of ROS and their derivatives are required for proper maintenance, proliferation, differentiation and apoptosis of stem cells and their committed progenitors (Kops *et al*, 2002; Giorgio *et al*, 2005; Tothova *et al*, 2007; Owusu-Ansah & Banerjee, 2009; Sinenko *et al*, 2010). In *Drosophila*, developmentally regulated levels of ROS are critical for maintenance of haematopoietic progenitors within the MZ of

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the lymph gland (Owusu-Ansah & Banerjee, 2009). In contrast, under normal growth conditions, PSC cells in wild-type larvae had very low levels of ROS expression (Fig 1A,A') compared with that in the progenitor population of cells within the MZ (Fig 1A). To induce oxidative stress in the PSC we inactivated ND75, a component of complex I of the electron transport chain (ETC; Owusu-Ansah & Banerjee, 2009), with double-stranded RNA (dsRNA) using the *Gal4/UAS* misexpression system (Brand & Perrimon, 1993) and the PSC-specific *Antp-Gal4* driver. As previously observed in other tissues (Owusu-Ansah et al, 2008), ND75 inactivation caused a readily detectable increase in ROS in the PSC cells (Fig 1B,B'), rising to levels similar to those seen in the progenitor cells of the MZ. The phenotypic consequence of inducing oxidative stress in the cells of the PSC was a remarkably robust increase in numbers of circulating lamellocytes (Fig 1C–E). Such an elevated number of lamellocytes was usually observed in wild-type larvae only if they were infested by parasitic wasps (Fig 2E,H). Although *Antp-Gal4* is not expressed anywhere in the blood system, except the PSC, this driver is also expressed in other larval tissues. To exclude the possibility that the effect was due to a non-PSC expression of *Antp-Gal4*, the function of ND75 was also eliminated using the *Dot-Gal4* driver normally expressed at high levels in the PSC, and this resulted in an identical lamellocyte response (Fig 1C). In contrast, oxidative challenge to various other larval tissues, including the fat body (*LSP2-Gal4*), the epidermis (*A58-Gal4*), the neurons (*C127-Gal4*), the dorsal vessel (*Hand-Gal4*), the ring gland (*5015-Gal4*), the wing imaginal disc (*ap-Gal4*) or the trachea (*btl-Gal4*), did not have a significant effect on lamellocyte differentiation (supplementary Fig S1A online). Furthermore, high ROS levels generated within the progenitor cells (*dome-Gal4*) of the lymph gland, which causes autonomous differentiation of this population (Owusu-Ansah and Banerjee, 2009), also did not have any significant effect on the non-autonomous differentiation of lamellocytes in the circulation (supplementary Fig S1A online). In contrast, oxidative challenge of the PSC caused non-autonomous lamellocyte response in circulation as well as within the lymph gland (Fig 1D'–E'). The PSC-mediated effect was due to mitochondrial dysfunction and not specifically linked to the product of the ND75 gene, because attenuation of *PDSW* (another complex I component), cytochrome-*c* oxidase, subunit Va (*CoVa*, a component of ETC complex IV) or *Marf* (mitochondrial assembly regulatory factor) function in the PSC, all induced increases in lamellocyte differentiation (supplementary Fig S1B online). The strength of the lamellocyte response to complex I inactivation depended on the strength of the dsRNA construct used in the experiment. Temporally, induction of the mutation in the second-larval instar caused the lamellocyte response to be seen in the third instar (not shown). This correlates well with the timescale of response to parasitic wasp infection (Croizatier et al, 2004). Finally, this oxidative stress elicited a cell-specific response; for example, no significant effect was seen on the differentiation of crystal cells and plasmatocytes in circulation (supplementary Fig S1C,D online). These results establish that the oxidative status of the PSC has a specific and non-autonomous role in lamellocyte differentiation as an immune response to parasitic invasion.

Akt1/FoxO controls oxidative sensing of the PSC niche

The status of the PSC cells on oxidative stress conditions was further analysed in some detail. ND75 dysfunction does not affect

proliferation or maintenance of the PSC, because the number of PSC cells, which maintain expression of *Antp*, remains intact in this mutant background (Fig 1A–B'). In addition, no apoptosis is detected in ND75-deficient PSC cells (not shown; Owusu-Ansah et al, 2008; Owusu-Ansah & Banerjee, 2009), and also, apoptosis in the PSC alone, specifically induced by overexpression of *Hid/Rpr*, has no effect on lamellocyte differentiation (supplementary Fig S1E online).

Overexpression of superoxide dismutase-2 (SOD2) as a scavenger for ROS (Finkel & Holbrook, 2000) in ND75-deficient PSC is able to suppress the lamellocyte response significantly (Fig 1F). Furthermore, activation of the Forkhead box O (FoxO) transcription factor that positively regulates expression of anti-oxidant enzymes, including SOD2 (Nemoto & Finkel, 2002), completely suppresses the *dsND75*-induced lamellocyte response (Fig 1G). Inactivation of the Akt1 protein kinase in PSC also results in a near-complete suppression of the *dsND75*-induced lamellocyte response (Fig 1G), suggesting a role for the PI3K/Akt pathway in the regulation of FoxO. This is an important issue because FoxO activity can also be controlled by the Jun N-terminal kinase (JNK) pathway (Essers et al, 2004; Owusu-Ansah & Banerjee, 2009), but in the PSC the AKT pathway mediates this effect. The JNK reporter (*puc69-lacZ*) is not expressed in the PSC, and inactivation of JNK (encoded by the *basket* gene) using the dominant-negative form (*bsk^{DN}*) does not suppress *dsND75*-induced lamellocyte response (supplementary Fig S2A online). The FoxO reporter (*4E-BP-lacZ*) is robustly activated in the ND75-deficient PSC (supplementary Fig S2B–C' online); however, loss of translational inhibition mediated by 4E-BP does not mimic this effect (not shown). It is important to point out that under wild-type non-stressed conditions, the PSC has relatively low levels of ROS, and therefore inactivation of either *Foxo* or *SOD2* has no phenotypic consequence (supplementary Fig S2D online). We interpret these data to indicate that metabolic dysfunction induces an oxidatively stressed PSC that causes the activation of this pathway and the lamellocyte response.

Spitz from the PSC activates cellular immune response

Differentiation of lamellocytes has been associated with the JAK/STAT, JNK and Ras/Erk signalling pathways (Harrison et al, 1995; Sorrentino et al, 2004; Zettervall et al, 2004; Williams et al, 2006; Makki et al, 2010). We genetically altered these pathways in an ND75-deficient PSC background to identify which, if any, is involved in the lamellocyte response. Inactivation of the unpaired ligands (*upd3*, *upd2* or *upd*) that activate the JAK/STAT pathway or of *eiger* (*egr*), which activates JNK signalling, did not suppress the lamellocyte phenotype (supplementary Fig S3A online). This strongly suggests that these pathways are not involved in the process downstream of ROS in the PSC and is consistent with previous studies showing that components of the JAK/STAT pathway (*upd3*, *dome* and *Tep4*; Krzemien et al, 2007; Makki et al, 2010) and JNK (*puc69-lacZ* reporter) are not involved in the functioning of the PSC. However, these pathways are likely to be involved in direct regulation of lamellocyte differentiation independently of the PSC function (Zettervall et al, 2004; Makki et al, 2010). In contrast, inactivation of *spitz* (*spi*), encoding the ligand for epidermal growth factor receptor (EGFR), in the context of ND75-deficient PSC significantly suppresses the lamellocyte response (Fig 2A). Furthermore, overexpression of the secreted

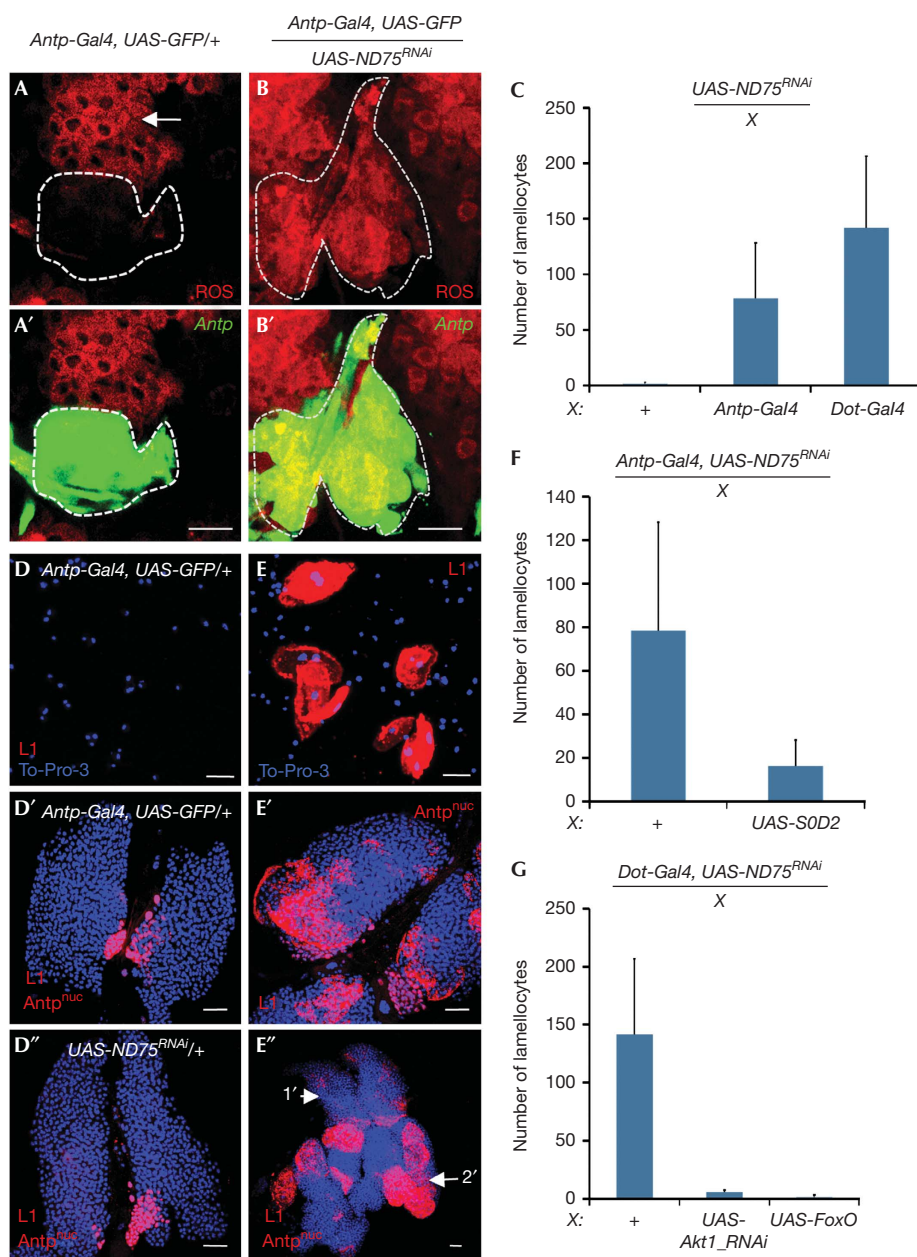
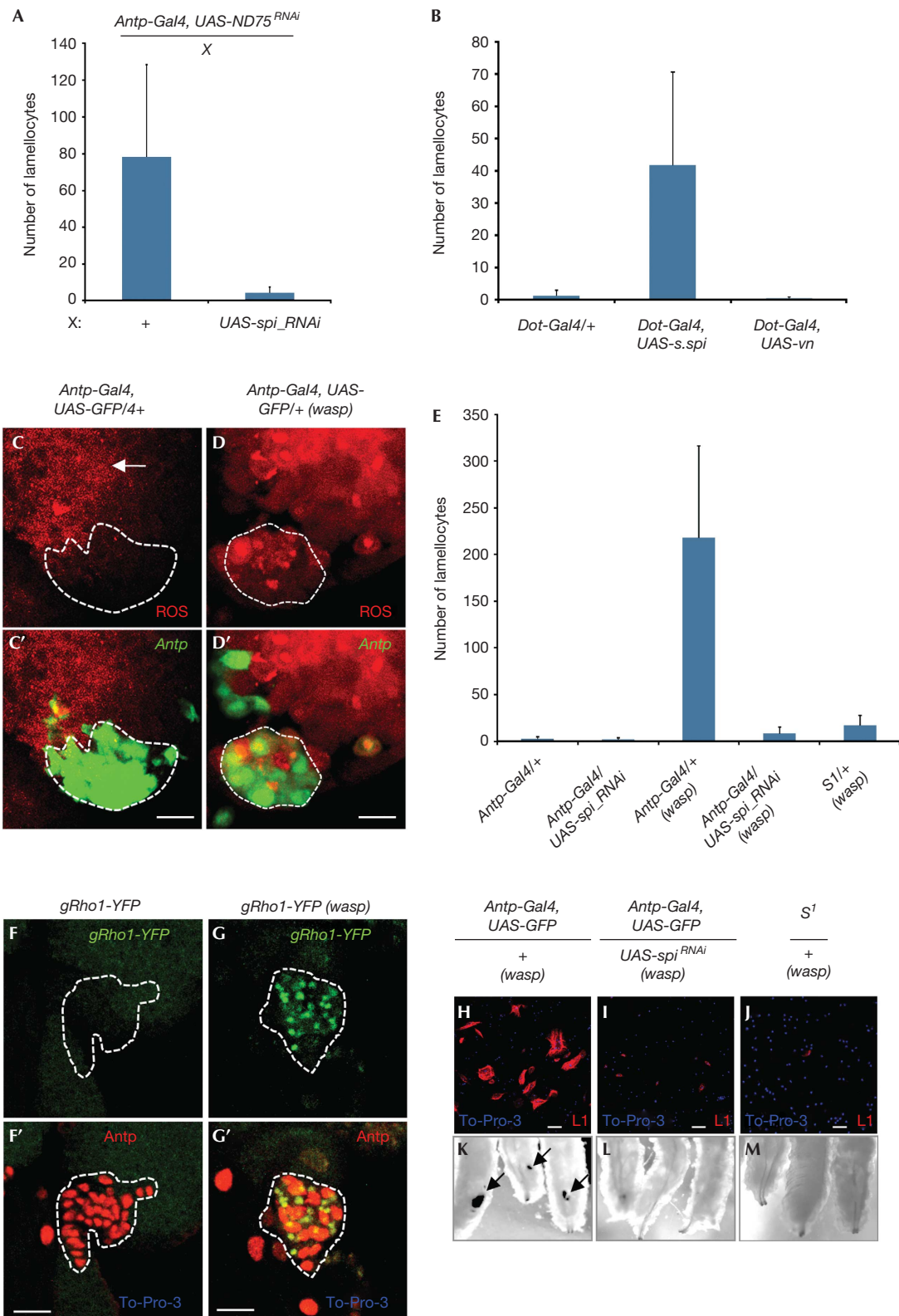


Fig 1 | ETC-mediated oxidative stress in the posterior signalling centre induces differentiation of lamellocytes. Cell markers are colour coded and indicated on each panel; genotypes are indicated at the top of each panel. All error bars represent standard deviation from the mean. (A,A') Posterior signalling centre (PSC) cells (marked by *Antp-Gal4, UAS-GFP*, green, outlined) have very low levels of reactive oxygen species (ROS) compared with neighbouring progenitor population (arrow, red). (B,B') Inactivation of *ND75* in the PSC (green) causes significant upregulation of ROS expression in these cells (red). Scale bar, 10 μ m. (C) Lamellocytes are rarely, if at all, seen in circulation of the wild-type larvae (+); $n = 14$. Oxidative stress induced by inactivation of *ND75* (*UAS-ND75^{RNAi}*) in the PSC cells with the use of *Antp-Gal4* ($n = 14$) or *Dot-Gal4* ($n = 16$) causes a significant increase in the number of circulating lamellocytes. Additional drivers are included as *Gal4* lines shown along the X axis. (D-D'') In wild-type larvae, lamellocytes (L1, red membrane stain) do not develop in circulation (D) or within the lymph gland (D',D''); genotypes indicated; scale bar, 25 μ m). The red staining in the PSC region is nuclear, marks *Antp* and is distinguishable from the extensive L1 membrane staining (as in E-E''). (E-E'') Inactivation of *ND75* in PSC causes extensive lamellocyte differentiation in larval circulation (E), the primary (E') and secondary (arrow, E'') lobes of the linkage group. (F,G) Superoxide dismutase-2 (SOD2) and Forkhead box O (FoxO) are crucial in controlling lamellocyte response in the PSC. The controls (+), show that inactivation of *ND75* (*UAS-ND75^{RNAi}*) in the PSC with *Antp-Gal4* (F, $n = 14$) or *Dot-Gal4* (G, $n = 14$) causes significant increase in lamellocyte numbers. Overexpression of SOD2 (F, $P < 0.001$, $n = 18$) or Foxo (G, $n = 14$) or inactivation of *Akt1* (G, $n = 14$) in the *ND75*-deficient PSC suppresses this lamellocyte response. Misexpression constructs are indicated along the X axis. ETC, Electron transport chain; GFP, green fluorescent protein; RNAi, RNA-mediated interference; UAS, upstream activating sequence.



form of Spi (s.Spi), but not the alternative EGFR ligand, Vein (Vn; Golembo *et al*, 1996; Schnepf *et al*, 1996) in the PSC, causes increased differentiation of circulating lamellocytes (Fig 2B) in an

otherwise wild-type larva. EGFR mutant *Egfr^{TS}/Egfr¹⁸* lymph glands develop normally (supplementary Fig S3B,C online), suggesting that EGFR signalling is not required for normal lymph gland

◀ **Fig 2** | Spitz signal from the oxidatively challenged PSC is required for the development of the lamellocyte response. (A) Inactivation of *ND75* in PSC causes a significant increase in numbers of circulating lamellocytes (+); $n = 14$. Inactivation of *spitz* (*spi^{RNAi}*) in the *ND75*-deficient PSC significantly suppresses the lamellocyte numbers; $n = 16$. (B) Overexpression of secreted Spitz (*Dot-Gal4*, *UAS-s.spi*, $n = 11$) but not the alternative epidermal growth factor receptor (EGFR) ligand *vein* (*Dot-Gal4*, *UAS-vn*, $n = 10$) in the PSC causes a significant increase in the lamellocyte numbers, compared with the wild-type background (*Dot-Gal4/+*, $n = 14$). (C,C') Under normal growth conditions, PSC cells (marked by *Antp-Gal4*, *UAS-GFP*, green outlined) are largely negative for reactive oxygen species (ROS; red); scale bar, 10 μm . (D,D') ROS levels (red) are significantly increased in the PSC cells after 12 h of wasp infestation. (E) Lamellocytes do not develop in wild-type (*Antp/+*, $n = 12$) background or when *spitz* (*spi*) is inactivated in the PSC with the use of *Antp-Gal4* (*Antp-Gal4*, *UAS-spi^{RNAi}*, $n = 15$). Large numbers of lamellocytes develop in wasp-infested wild-type larvae (*Antp/+* (*wasp*), $n = 10$). Loss of *spi* function in the PSC (*Antp-Gal4*, *UAS-spi^{RNAi}*(*wasp*), $n = 15$) or loss of a single copy of *Star* (*S¹/+*, $n = 15$) results in a suppression of lamellocyte production in the infested mutant larvae. (F,F') Under normal growth conditions PSC cells (marked by *Antp*, red outlined) do not express Rho (*gRho1-YFP*, green); scale bar, 10 μm . (G,G') Rho (*gRho1-YFP*, green) is expressed as punctate dots in the cells of the PSC after 12–18 h of wasp infestation. (H) Large numbers of circulating lamellocytes develop in larvae infested with wasp eggs; scale bar, 50 μm . (I) Differentiated lamellocytes do not develop in the wasp-infested larvae that lack *spi* function in the PSC. (J) Differentiated lamellocytes do not develop in wasp-infested larvae that lack of single dose of S protein. (K) Melanotic capsules (arrows) are seen in wasp-infested larvae. (L) Melanotic capsules do not develop in the wasp-infested larvae that lack *spi* function in the PSC. (M) Melanotic capsules do not develop in the wasp-infested larvae of mutants lacking a single copy of *Star*. Cell markers are colour coded and indicated on each panel; genotypes are indicated at the top of each panel. The error bars shown in A, B and E represent standard deviation from the mean. GFP, green fluorescent protein; PSC, posterior signalling centre; RNAi, RNA-mediated interference; UAS, upstream activating sequence; vn, Vein; YFP, yellow fluorescent protein.

development but rather is involved in the regulation of a cellular immune response as a signalling event from the PSC only when the latter is oxidatively stressed.

The PSC-dependent parasitic challenge induced by wasp egg infestation and the mechanism described above both give rise to the same cellular response. We therefore decided to determine whether parasitization causes oxidative stress to the PSC. We found that immune challenge caused by wasp infestation induces high levels of ROS in the PSC cells as seen 12 h after invasion (Fig 2C–D'). The most prominent effect is on superoxide radicals detected with dihydroethidium staining (Fig 2D,D'); a smaller but detectable elevation of peroxide radicals revealed by RedoxSensor staining (supplementary Fig S4A–B' online) is also apparent in PSC cells on this immune challenge. Scavenging these ROS types in the PSC by overexpressing SOD2 or catalase (Cat; Finkel & Holbrook, 2000) but not glutathione peroxidase (GPx), which reduces thioredoxin-mediated effects (Missirlis et al, 2003), significantly suppresses the lamellocyte response caused by wasp infestation (supplementary Fig S4C online). These genetic results are consistent with a model in which parasitic infection by wasp eggs raises ROS levels in the PSC, which then causes lamellocyte induction by expressing Spitz (Fig 3E). To test this model, we inactivated *spi* within the PSC in larvae infected by parasitic wasps. This caused a strong suppression of the lamellocyte response (Fig 2E,H,I,K,L); the few remaining L1 marker-positive cells are immature, as indicated by their relatively small cell size and their morphology (Fig 2H,I). In addition, melanotic capsules that are indicative of extensive cellular immune response to parasitic infection do not develop in a *spi* mutant background during wasp infestation (Fig 2K,L). Inactivation of *spitz* in the PSC did not affect the increase in ROS triggered by wasp infestation (supplementary Fig S4D–F' online). Thus *spi* does not regulate the ROS levels in the PSC; rather, wasp infection raises ROS levels, which leads to release of the s.Spi. Previous studies have shown that s.Spi production requires the function of the trafficking protein Star (S), and the protease Rhomboid (Rho1; Kolodkin et al, 1994; Urban et al, 2001; Tsruya et al, 2002). We found that the wasp-induced lamellocyte response and melanotic capsule

formation are robustly suppressed on the loss of a single copy of *Star* (Fig 2E,J,M). More importantly, parasite-induced immune challenge specifically upregulates Rho1 in the PSC (Fig 2F,G') by an as yet unidentified mechanism. These data establish that S and *Rho1* are canonically required for processing and releasing the Spitz from the PSC.

Secreted Spitz is known to bind to EGFR and activate the Ras/Erk pathway (Seger & Krebs, 1995). A dominant-negative form of EGFR (*Egfr^{DN}*) strongly suppresses the lamellocyte response induced by wasp infestation when it is expressed in the lymph gland and the circulating haemocytes using the pan-haemocyte *HHLT Gal4* driver (Fig 3A). This phenotype is virtually identical to that seen when *spi^{RNAi}* is expressed in the PSC using *Antp-Gal4*. In addition, we used compartment-specific drivers and found that inactivation of the receptor in the cortical zone of the lymph gland and in circulating haemocytes (using lineage-traced *Hml^Δ-Gal4* line) prevents *Hml*-positive cells from becoming lamellocytes on wasp infestation (supplementary Fig S4G online). Importantly, we also found that a small subset of lamellocytes does not express *Hml* in the wild-type background and consequently *Egfr^{DN}* is not expressed in these cells when *Hml^Δ-Gal4* is used as a driver. These *Hml⁻*, L1+ lamellocytes are easily detectable in this genetic background and act as an internal control. Expression of an activated form of EGFR (*Egfr^{Act}*) in *Hml⁺* haemocytes causes a robust increase in lamellocyte differentiation (Fig 3A). This is also consistent with previous work, which showed that activated Ras induces an increase in the total number of haemocytes, including lamellocytes (Asha et al, 2003). Finally, both loss of *ND75* in the PSC and wasp infestation cause robust activation of Erk as evident by an increase in dpErk staining in circulating haemocytes including lamellocytes (Fig 3B–D). This indicates that lamellocytes in circulation differentiate from precursor cells on activation of Spi/EGFR/Erk signalling. Taken together, these data provide strong support for the model shown in Fig 3E.

PSC cells have two independent functions: they serve as a haematopoietic niche in the lymph gland, where they orchestrate the maintenance and proper differentiation of haematopoietic progenitors (Krzemien et al, 2007; Mandal et al, 2007), and they

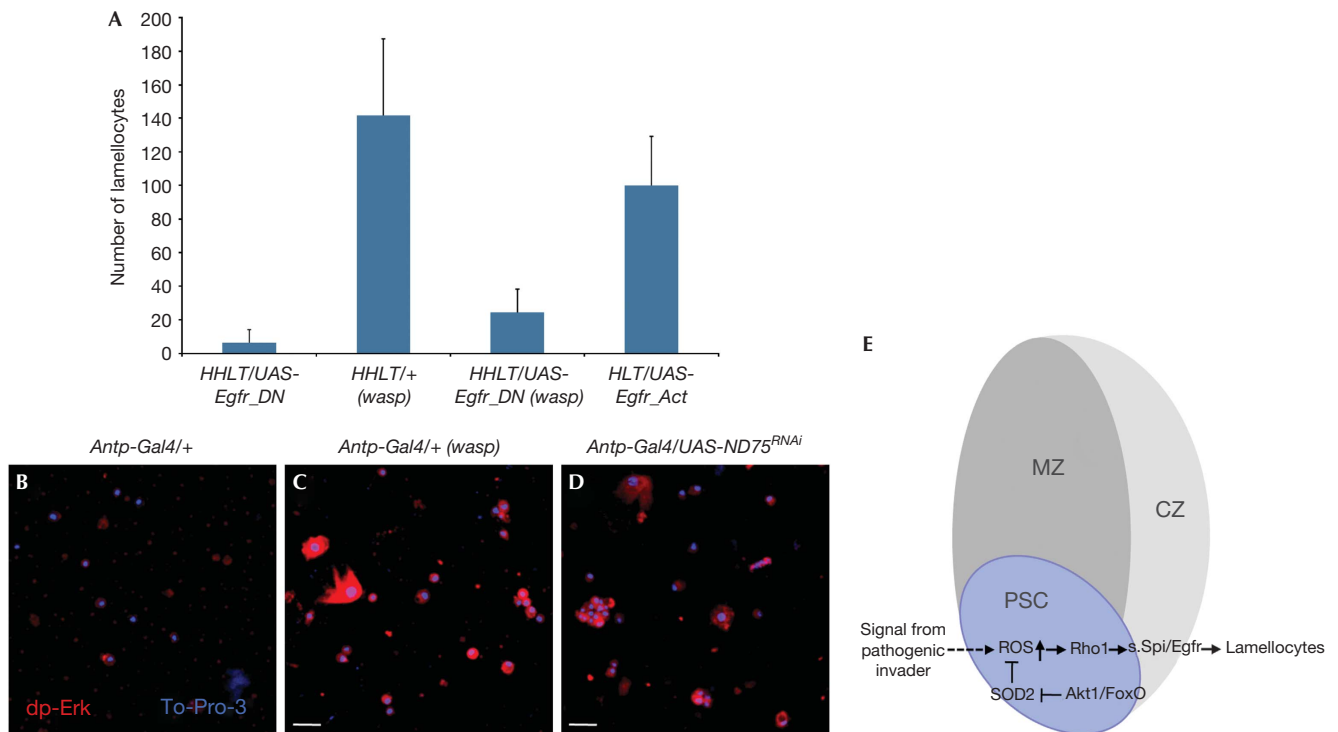


Fig 3 | Spi/EGFR signalling mediates lamellocyte cellular immune response. (A) As controls, lamellocytes are barely detected in wild-type circulation or when *Egfr* is inactivated in all haemocytes (*HHLT-Gal4*, *UAS-Egfr^{DN}*, $n = 10$). All error bars shown represent standard deviation from the mean. Large numbers of lamellocytes develop in wasp-infested wild-type larvae (*HHLT-Gal4/+ (wasp)*, $n = 10$). Loss of *Egfr* function in all haemocyte populations (*HHLT-Gal4*, *UAS-Egfr^{DN}(wasp)*, $P = 0.0001$, $n = 10$) results in a suppression of lamellocyte production in the infested mutant larvae. Overexpression of *Egfr^{Act}* in the haemocyte precursors (*HLT-Gal4*, *UAS-Egfr^{Act}*) induces the generation of large number of lamellocytes. (B–D) Under normal growth conditions dpErk expression is extremely low (B, red). Oxidative stress in the PSC induced by wasp infestation (C, red) or by inactivation of ND75 in the PSC (D, red) causes a significant increase in dpErk-positive cells in larval circulation. Genotypes are indicated at the top of the panels. Scale bar, 25 μm . (E) Proposed model. Signal from parasitic invader induces changes in the oxidative status of the PSC that in turn causes the production of secreted Spitz (s.Spi) protein, which signals via EGFR receptor, inducing lamellocyte differentiation. The Akt/FoxO pathway is involved in the regulation of the oxidative status of the PSC. CZ, cortical zone; DN, dominant-negative; EGFR, epidermal growth factor receptor; FoxO, Forkhead box O; HHLT, *Hand-Hml* lineage traced; HLT, *Hml* lineage traced; MZ, medullary zone; PSC, posterior signalling centre; ROS, reactive oxygen species; SOD, Superoxide dismutase-2; *Spi*, *Spitz*.

regulate the cellular immune response by controlling lamellocyte differentiation in response to infection (Crozatier *et al*, 2004; Makki *et al*, 2010). The results presented here establish the mechanism for this latter function. Changes in oxidative status, caused by events of parasite invasion or ETC dysfunction, initiates a signal within this immunocompetent compartment causing the secretion of a cytokine ligand, Spitz, that induces differentiation of lamellocyte precursors in the circulatory system of the larva (Fig 3E). The identified mechanism is consistent with previously reported studies in mammals, which have shown that mitochondrial ROS can trigger systemic signals that reinforce the innate immune response (Arsenijevic *et al*, 2000). Our studies raise the possibility that specific populations of cells also exist in mammalian systems that sense oxidative stress due to infection and non-autonomously signal myeloid progenitors to initiate differentiation and enhance the immune response. Whether such populations are to be found within the haematopoietic niche as in *Drosophila* remains a speculation that can be tested in future studies.

METHODS

For sources of stocks and antibodies used, please see supplementary information.

Genetics. *Hand-Hml* lineage-traced Gal4 (*HHLT Gal4*) and *Hml* lineage-traced Gal4 (*HLT Gal4*) lines were used to express transgenes in all haemocyte populations and *Hml*-positive haemocytes, respectively (Evans *et al*, 2009); Evans and Banerjee, unpublished). Two independent insertions of each UAS-dsRNA constructs were used. For controls, larvae from crosses of particular *Gal4* line and w^{1118} were used. *Egfr^{TS}/Egfr¹⁸* mutants were shifted to restrictive temperature (29 °C) during the late embryonic stage—early first-larval instar and lymph glands analysed were dissected from third-instar larvae. Crosses were maintained on standard *Drosophila* cornmeal/sucrose/yeast medium at 25 or 29 °C.

Wasp infestation experiment. Late second or early third-instar larvae were treated with wasps (*Leptopilina bouleardii*) for 12 h at room temperature. At 12 or 48 h (25 °C) after infestation (Crozatier *et al*, 2004), haemocyte samples were collected from

third-instar larvae on glass slides and immunostained with L1 antibodies as described (Sinenko *et al*, 2004). Statistical analyses were performed using Student's *t*-test. Data in the figures represent mean \pm s.d.

Supplementary information is available at EMBO reports online (<http://www.emboreports.org>).

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Author contributions: S.A.S. and U.B. developed the project. S.A.S. designed and performed experiments. J.S. performed experiments. S.A.S. and U.B. discussed results and wrote the manuscript.

CONFLICT OF INTEREST

The authors declare that they have no conflict of interest.

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