INTRODUCTION

Occlusive arterial disease is the narrowing of an artery, resulting in impaired blood flow and ischaemia distal to the obstruction. To prevent or resolve ischaemia, neovascularization is induced spontaneously to restore blood flow. Angiogenesis and arteriogenesis are the main components of neovascularization and are induced under pathological conditions, such as peripheral arterial disease and myocardial infarction.

Angiogenesis, the sprouting of new capillaries from pre-existing vasculature, responds to hypoxia by up-regulating pro-angiogenic growth factors such as vascular endothelial growth factor (VEGF).
These growth factors induce endothelial cells from the pre-existing vasculature to migrate, grow and differentiate to shape new capillaries.\textsuperscript{3,4} Arteriogenesis, the maturation of pre-existing collateral vessels into arterioles,\textsuperscript{5,6} is induced by inflammation, shear stress and circumferential stretch on the vascular wall after occlusion, mainly proximal to the occlusion. Inflammatory cells such as macrophages, monocytes, but also CD4+ and CD8+ T cells adhere and invade into the vascular wall and produce inflammatory cytokines, chemokines and growth factors, which cause further maturation of arterioles and induce angiogenesis. Subsequently, smooth muscle cells proliferate and migrate towards the vessel wall, resulting in full functional arterioles.\textsuperscript{7,9,10} Both angiogenesis and arteriogenesis are essential for accurate neoangiogenesis, to restore blood flow after arterial occlusions.

For the production of inflammatory cytokotyks, chemokines and growth factors, inflammatory cells require activation. Substantial activators are pattern recognition receptors, such as toll-like receptors (TLR).\textsuperscript{11,12} TLR3 expressed in intracellular vesicles recognizes dsRNA, which activates a Toll/interleukin-1 receptor domain-containing adapter-inducing interferon-β protein (TRIF)-dependent pathway. The TLR3-TRIF pathway induces type-I IFN production, via interferon regulatory factor (IRF)3 and IRF7 phosphorylation, and pro-inflammatory cytokine production via transcription factors nuclear factor kappa-light-chain-enhancer of activated B cells (NFκB).\textsuperscript{12-15} IRF3 and IRF7 are mainly phosphorylated downstream TLR3, but might also be partly activated via other TLRs (e.g TLR4, TLR7, TLR9).\textsuperscript{12,15} It is assumed that NFκB activation results in pro-inflammatory cytokine production and IRF3/IRF7 phosphorylation and results in type-I IFN production. However, it is observed that TLR3 can also activate pro-inflammatory transcription factors via IRF3 and IRF7 and may be potential regulators of cell proliferation and survival.\textsuperscript{16-19} Since neoangiogenesis is mainly induced by pro-inflammatory cytokines and growth factors, we hypothesize an important role for IRF3 and IRF7 in arteriogenesis and angiogenesis.

In the current study, we aimed to elucidate the role of IRF3 and IRF7 in vivo in post-ischaemic neoangiogenesis in a hind limb ischaemia (HLI) mouse model, using wild-type C57BL/6, Irf3\textsuperscript{−/−} and Irf7\textsuperscript{−/−} mice.

\section{Material and Methods}

\subsection{Mice}

This study was performed in compliance with the Dutch government guidelines and the Directive 2010/63/EU of the European Parliament. All experiments were approved by the committee on animal welfare of the Leiden University Medical Center (Leiden, the Netherlands). For the experiments 10-18-week-old male mice were used. C57BL/6 mice were purchased from Charles River Laboratories and Irf3\textsuperscript{−/−} and Irf7\textsuperscript{−/−} mice were kindly provided by Dr Taniguchi (University of Tokyo, Japan) and bred in our facility.\textsuperscript{20} Mice were fed a chow diet ad libitum.

\subsection{Murine HLI model}

Mice were anaesthetized with an intraperitoneal injection of midazolam (8 mg/kg; Roche Diagnostics), medetomidine (0.4 mg/kg; Orion), and fentanyl (0.08 mg/kg; Janssen Pharmaceuticals). Unilateral hind limb ischaemia (HLI) was performed. In brief, a single ligation model was performed by electrocoagulation of the left common femoral artery proximal to the bifurcation of the popliteal and saphenous artery.\textsuperscript{21,22} After surgery, anaesthesia was antagonized with flumazenil (0.7 mg/kg, Fresenius Kabig). Buprenorphine (0.1 mg/kg, MSD Animal Health) was given after surgery to relieve pain in a fixed regime and when circumstances required this was repeated. Number of total operated mice was 11 per group, however, in the C57BL/6 group 5 mice deceased during or after surgery and 2 and 4 mice, respectively in the Irf3\textsuperscript{−/−} and Irf7\textsuperscript{−/−} group. Mice characteristics of Irf3\textsuperscript{−/−}, Irf7\textsuperscript{−/−} and C57BL/6 mice are shown in Table S1. Body weight at day of surgery (t = 0) and when killed (t = 28) was similar in the Irf3\textsuperscript{−/−} and Irf7\textsuperscript{−/−} mice compared to C57BL/6 mice.

\subsection{Laser Doppler perfusion imaging}

Post-ischaemic blood flow recovery was measured in the left ischaemic and right non-ischaemic paw of Irf3\textsuperscript{−/−}, Irf7\textsuperscript{−/−} and C57BL/6 mice with the use of laser Doppler perfusion imaging (LDPI) (Moor Instruments).\textsuperscript{23} Blood flow was measured before and directly after surgery and at day 3, 6, 9, 14, 21 and 28. Paw perfusion was expressed as a ratio of left ischaemia to right non-ischaemic paw.\textsuperscript{24} Before LDPI, mice were anaesthetized as described before\textsuperscript{25,26} with an intraperitoneal injection of midazolam (8 mg/kg) and medetomidine (0.4 mg/kg). After LDPI, anaesthesia was antagonized by subcutaneous injection of flumazenil (0.7 mg/kg). Mice were killed 28 days after surgery after the last LDPI via cervical dislocation after subcutaneous admission of analgesic fentanyl (0.08 mg/kg). The adductor, soleus and gastrocnemius muscles were harvested for either (immuno) histochemical analysis or RT-qPCR.

\subsection{Immunohistochemistry}

Adductor muscles were fixed in 4% formaldehyde and embedded in paraffin, soleus and gastrocnemius muscles were snap frozen on dry ice and stored at −80°C. Cross sections of 6 μm were made throughout the embedded adductor muscle. Adductor muscle sections were stained with alpha smooth muscle cell actin (aSMActin, DAKO) to visualize vascular smooth muscle cells (VSMC). Stained slides were photographed (20× magnification) with microscope photography software (Axiovision, Zeiss) and analysed with ImageJ (FIJI) by counting the number of arterioles and measuring the diameters of each arteriole with a visible lumen to determine arteriogenesis.\textsuperscript{25} In addition, adductor muscle sections were stained with alpha smooth muscle cell actin (pink) and MAC3 (green) (BD Pharamingen) to visualize macrophages around arterioles. Images were acquired on a Philips Ultra Fast Slide Scanner (Philips Digital Pathology Solutions, Best) and analysed by counting the number of
macrophages around the arterioles divided by the circumference of the arteriole (presented as macrophages/μm). However, 12 sections, in the same area in all mice were analysed per mice per leg and the average was used for analyses. Negative controls were performed by omitting either the secondary antibody of aSMActin or MAC3, or PBS.

Fresh-frozen soleus muscles were cross sectioned in 6 μm slices with a cryostat and sections were fixed in acetone. To visualize endothelial cells, soleus muscle sections were stained with CD31 (BD Pharmingen). Stained slides were photographed (20× magnification) with microscope photography software. The number of CD31+ structures was analysed with ImageJ, which was used to determine angiogenesis and was presented in absolute numbers of CD31+ structures per area.

2.5 | RNA isolation, cDNA synthesis and RT-PCR from adductor muscles

RNA was isolated from snap frozen gastrocnemius muscles 28 days after surgery as described before. In brief, muscle tissues were crushed with mortar and pestle, while using liquid nitrogen to preserve sample integrity. RNA was isolated according to manufacturer’s protocol using TRIzol Reagent (Life Technologies) (FFPE RNA isolation kit, Qiagen), after which sample concentration and purity were examined by nanodrop (Nanodrop Technologies). Complementary DNA (cDNA) was synthesized using a High-Capacity cDNA Reverse Transcription Kit according to the manufacturer’s protocol (Applied Biosystems). To measure the expression of *tnfa* (forward TGAACCTCGGTTGATCGG, reverse CTCCCTACCTTGGTTTGTG), *ccl2* (forward TATGCTGACCTGCTGTGTGAT, il6 (forward TCGGAGAAGAGACTTCACA, reverseTGGCATTTGGCAC AAACCTTT), il1β (forward ACGTCTCGCTTGGCTTTC, reverse TGGACTGGAAGCCTGGATTG), *sdf1* (forward TGGGCTTCCGAACCTTAAAT, reverse AGCAGAGCCCTGGAATGTG), *vegf164* (forward AGCTGCTGAGGACTTGCTTC, reverse ATGAGTCCGCTTCCG), *vegfr1* (forward ACGGAGGACCTGGCTTTC, reverse AGCTGCATGTCGACTTAC), *vegfr2* (forward CGTGCAGAGGACCTGGCTTTC, reverse AGCTGCTGAGGACTTGCTTC), RT-qPCR was performed using QuantiTect SYBR Green PCR Kit (Qiagen). GAPDH (Applied Biosystems) was used as a housekeeping gene. All RT-qPCRs were performed on a 7500/7500 Fast Real-Time PCR System (Applied Biosystems) and the 2−ΔΔCt method was used to analyse the relative changes in gene expression.

2.6 | Statistical analysis

All data are presented as mean ± SEM. In statistics software GraphPad Prism 7.0, statistical analyses on parametric data were performed by using a 2-tailed Student’s t test to compare individual groups. Mann-Whitney test was used for nonparametric data. A 1-way ANOVA was performed on parametric data comparing more than two groups and a Kruskal-Wallis test was performed on non-parametric data, followed by Tukey’s post hoc test. P value of <0.05 was considered significant.
Irf7−/− mice and analysed paw perfusion before ligation and 3, 6, 9, 14, 21 and 28 days after ligation. Results of the LDPI, expressed as the left ischaemic/right non‐ischaemic ratio, are shown in Figure 1. After ligation, the paw perfusion decreased tremendously in all groups to ratios of approximately 0.05. The fact that these ratios are similar in all groups, suggests that the number of pre‐existing collaterals, that allow residual blood flow to the paw after ligation, is similar. C57BL/6 mice already recovered almost 50% 3 days after surgery, whereas the Irf3−/− (P = 0.0005) and Irf7−/− (P = 0.01) mice only recovered 20%. Both Irf3−/− (P = 0.006) and Irf7−/− (P = 0.006) mice also showed an impaired blood flow recovery compared to C57BL/6 mice 6 days after ligation. C57BL/6 mice already reached their maximal recovery 6 days after surgery, showing a similar paw perfusion as before the ligation, which did not change until sacrifice. After 9 days, Irf3−/− and Irf7−/− mice reached their maximal recovery of, respectively, 78% and 75% which did not further increase up to the time of sacrifice, 4 weeks after surgery. Both Irf3−/− and Irf7−/− mice did not manage to reach the same level of post‐ischaemic blood flow recovery as C57BL/6 mice. This indicates that IRF3 and IRF7 have an important role in post‐ischaemic blood flow recovery.

3.2 | Angiogenesis in soleus muscles

To investigate the underlying cause of the reduced post‐ischaemic blood flow recovery in Irf3−/− and Irf7−/− mice, we determined the angiogenic capillary formation in the soleus muscles of Irf3−/−, Irf7−/− and C57BL/6 mice. Soleus muscles of the left ischaemic and right non‐ischaemic calf were stained with CD31 (Figure 2A‐C). The number of CD31+ cells, expressed as left ischaemic/right non‐ischaemic, in the soleus muscle of Irf3−/− mice (P = 0.02) and Irf7−/− mice (P = 0.04) was decreased compared to C57BL/6 mice 28 days after surgery (Figure 2D). This suggests a decrease in angiogenesis. The number of CD31+ cells in the left ischaemic calf was significantly decreased in the soleus muscle of both Irf3−/− (P < 0.0001) and Irf7−/− (P = 0.003) mice compared to C57BL/6 mice (Figure 2E). The right non‐ischaemic calf showed no differences in number of CD31+ cells (Figure 2F) in Irf3−/− and Irf7−/− compared to C57BL/6 mice, indicating that there is no genetic difference in angiogenesis between Irf3−/−, Irf7−/− and C57BL/6 mice.

3.3 | Arteriogenesis in adductor muscles

In addition to angiogenesis, arteriogenesis also is an important process during neovascularization. To investigate arteriogenesis, adductor muscles were stained with aSMActin (Figure 3A‐C) and the diameter of arterioles were measured (Figure S1A,B) together with the number of arterioles. The diameter of the aSMActin+ arterioles, expressed as left ischaemic/right non‐ischaemic, was smaller in Irf3−/− mice (P = 0.04) and Irf7−/− mice (P = 0.02) compared to C57BL/6 mice 28 days after surgery (Figure 3D). The diameter of the aSMActin+ arterioles was decreased in the individual left ischaemic adductor muscles of Irf3−/− (P = 0.005) and Irf7−/− (P = 0.02) mice compared to C57BL/6 mice (Figure S1C). No significant differences were observed in the diameter of the aSMActin+ arterioles of the right non‐ischaemic adductor muscle of C57BL/6 compared to Irf3−/− and Irf7−/− mice.

**FIGURE 2** Angiogenesis in CD31+ stained soleus muscles. Representative image of the left ischaemic soleus muscle stained with CD31 (20× magnification) of (A) C57BL/6 mice (B) Irf3−/− mice and (C) Irf7−/− mice sacrificed 28 days after surgery (D) Ratio of the number of CD31+ cells in the soleus muscle of left ischaemic/right non‐ischaemic is shown. (E) The number of CD31+ cells in the soleus muscle of the left ischaemic calf is shown. (F) The number of CD31+ cells in the soleus muscle of the right non‐ischaemic calf is shown. Data is presented as mean SEM; *P < 0.05; **P < 0.01, ****P < 0.0001, a 2‐tailed Student’s t test was used. C57BL/6 n = 6, Irf3−/− n = 9, Irf7−/− n = 6
3.4 | Perivascular accumulation of inflammatory cells

Perivascular inflammation was demonstrated by the number of macrophages around the arterioles. An immunohistochemical double staining was performed on adductor muscles of Irf3−/− mice and C57BL/6 mice 28 days after HLI. Here, aSMActin was used to show the arterioles and MAC3 to show the macrophages around the arterioles. An immunohistochemical double staining was performed on adductor muscles of C57BL/6 mice killed 28 days after HLI. Here, aSMActin was used to show the arterioles and MAC3 to show the macrophages around the arterioles. An immunohistochemical double staining was performed on adductor muscles of C57BL/6 mice killed 28 days after HLI. Here, aSMActin was used to show the arterioles and MAC3 to show the macrophages around the arterioles.

3.5 | mRNA expression of essential genes for arteriogenesis and angiogenesis

RNA from the gastrocnemius muscles was isolated and the mRNA expression of tnfa, ccl2, il6, il1β, sdf1, vegf164, vegfr1 and vegfr2 was measured with qPCR as genes essential for angiogenesis and arteriogenesis. The relative mRNA expression of inflammatory cytokines tnfa, ccl2 and il6 was decreased in left ischaemic gastrocnemius muscles of Irf3−/− and Irf7−/− mice compared to C57BL/6 mice (Figure 5A-C). However, il1β and sdf1 were not differentially expressed in gastrocnemius muscles of Irf3−/−, Irf7−/− and C57BL/6 mice (Figure 5D,E). However, VEGF and the first receptor (VEGFR1) showed a trend towards a down-regulation in gastrocnemius muscles of Irf3−/− and Irf7−/− mice compared to C57BL/6 mice (Figure 5F,G). Interestingly, the mRNA expression levels of the second receptor (VEGFR2) were decreased in the left ischaemic gastrocnemius muscles of Irf3−/− and Irf7−/− mice compared to C57BL/6 mice (Figure 5H).

No differences were observed in the mRNA expression of tnfa, ccl2, il6, il1β, sdf1, vegf164, vegfr1 and vegfr2 in the right non-ischaemic gastrocnemius muscles of Irf3−/− and Irf7−/− mice compared to control mice (Figure S3). These results demonstrate that IRF3 and IRF7 regulate the expression of several genes essential for angiogenesis and arteriogenesis.

4 | DISCUSSION

In the present study, we observed a decreased post-ischaemic blood flow recovery in Irf3−/− and Irf7−/− mice compared to C57BL/6 mice, using a HLI model. This was supported by a decrease in angiogenesis and arteriogenesis in, respectively, soleus and adductor muscles of the Irf3−/− and Irf7−/− mice. Furthermore, mRNA expression levels of several pro-inflammatory cytokines (tnfa, il6, ccl2) and growth factor receptor (vegfr2) that are essential for the induction of angiogenesis and arteriogenesis, were decreased in the gastrocnemius muscles of Irf3−/− and Irf7−/− mice. This indicates a decreased post-ischaemic blood flow recovery caused by an impairment in angiogenesis and arteriogenesis due to lack of inflammatory components in the ischaemic tissue.
Angiogenesis provides a better distribution of blood to ischaemic tissue. Since we observed a decreased post-ischaemic blood flow recovery in the paw of Irf3−/− and Irf7−/− mice compared to C57BL/6 mice, we hypothesized attenuated angiogenesis in soleus muscles of Irf3−/− and Irf7−/− mice. This was confirmed when we measured a reduced angiogenic capillary formation in the soleus muscles of Irf3−/− and Irf7−/− mice compared to C57BL/6 mice. The reduced number of capillaries was only observed in the left ischaemic calf and not in the right non-ischaemic calf. This indicates that the genetic baseline in all mice was similar and the decreased ischaemia-induced angiogenesis in the ischaemic muscles of Irf3−/− and Irf7−/− mice was purely based on the effects of IRF3 and IRF7. Previously, a role of IRF3 and IRF7 in angiogenesis was indicated. A phenotypic genome-wide screening performed on human umbilical vein endothelial cell by Korherr et al identified factors that induce angiogenesis. Among these factors was the well-known factor VEGF as well as TRIF, kinase TBK1 and IRF3.28 This indicates that the TRIF/TBK/IRF3 pathway has angiogenic properties. In addition, IRF7 is also involved in angiogenesis since Jin et al also showed that IRF7 over-expression increased tumour formation with increased angiogenesis and cell heterogeneity by inducing inflammatory cytokine expression (IL6, CCL2 and CXCL1).29 This latter is in line with our results where we show an attenuated inflammatory response in absence of IRF7.

**FIGURE 4** Macrophages around arterioles. An immunohistochemical double staining was performed on adductor muscles of Irf3−/−, Irf7−/− and C57BL/6 mice sacrificed 28 days after HLI. aSMActin (pink) was used to show the arterioles and MAC3 (green) to show the macrophages around the collaterals. Representative images of the left ischaemic adductor muscle of (A) C57BL/6 mice, (B) Irf3−/− mice and (C) Irf7−/− mice stained with aSMActin and MAC3 is shown (scalebar = 100 μm) and a zoom in of the arterioles of the concerned (D) C57BL/6 mice, (E) Irf3−/− mice, (F) Irf7−/− mice arteriole is shown (scalebar = 10 μm). Arrows indicate where the macrophages are located in the arterioles. (G) Ratio (left ischemic/right non-ischemic) of the number of macrophages per μm circumference of arterioles in the adductor muscle is shown. (H) Number of macrophages per μm circumference of arterioles in the left ischemic adductor muscle is shown. (I) Number of macrophages per μm circumference of arterioles in the right non-ischemic adductor muscle is shown. Data is presented as mean SEM; **P < 0.01, Mann-Whitney test was used. C57BL/6 n = 5, Irf3−/− n = 7, Irf7−/− n = 7
Since angiogenesis is induced via VEGF, it is interesting that we found a down-regulation of VEGR2 in the ischaemic muscles of IRF3 and IRF7-deficient mice. VEGF can bind to VEGR1, VEGFR2 and VEGFR3, but VEGFR2 is the main receptor transmitting VEGF signals. The lack of VEGFR2 in the ischaemic tissue can (partly) explain the decrease in angiogenesis in IRF3 and IRF7-deficient mice compared to C57BL/6 mice. Interestingly, previously a relation between interferons, regulated by IRF3 and IRF7, and VEGF was observed. IFNs were able to mediate VEGF production from human mast cells and also in chronic myeloid leukaemia patients, imatinib in combination with pegylated-IFN-α2a was able to regulate VEGF levels.

Arteriogenesis is induced by inflammation, which results in the maturation of pre-existing collaterals into arterioles. For optimal neovascularization and thus post-ischaemic blood flow recovery, both angiogenesis and arteriogenesis are essential. We observed reduced arteriogenesis in the adductor muscles of Irf3−/− and Irf7−/− mice compared to C57BL/6 mice. The number of arterioles was only increased in Irf3−/− mice compared to C57BL/6 mice. However, the diameter of the arterioles was significantly decreased in Irf3−/− and Irf7−/− mice compared to C57BL/6 mice, which is characteristic for arteriogenesis since this is defined as the maturation of pre-existing collaterals. Interestingly, we showed a decrease in inflammatory cells around the arterioles and decreased mRNA expression of inflammatory cytokines IFNα, ccl2 and il6 in the gastrocnemius muscles. It is shown that several inflammatory cytokines and components are essential in the regulation of arteriogenesis such as TNFα, as well as the CCR7-CCL19/CCL21 Axis, NK cells and CD4+ T cells and CD27-CD70 T cell co-stimulation. We previously demonstrated that IRF3 regulates pro-inflammatory cytokines downstream TLR4 activation in vitro, in VSMCs and macrophages, prominent inflammatory cell types involved in arteriogenesis. Together with the diminished inflammatory response observed here, this partly explains how IRF3 and IRF7 affect neovascularization.

Although from the current data on cytokine expression and macrophage presence at 28 days, it cannot be excluded that one might also conclude that due to the reduced neovascularization response, inflammatory cytokine expression is reduced and not vice versa. Previous data from our and other groups have clearly shown that in early stage after induction of hind limb ischaemia hampered inflammatory responses lead to reduced neovascularization responses. The regulation of pro-inflammatory cytokines via IRF3 and IRF7, in general goes via the TRIF-NFκB pathway. TRIF is also involved in TLR3-mediated type-I IFN production via IRF3 and IRF7 activation. It is suggested that activation of TLR3-TRIF can also suppress angiogenesis via initiation of apoptotic cell death. The ability of TRIF to induce apoptosis however, was not dependent on its ability to activate either IRF3 or NFκB but was dependent on RIP, Fas-associated death domain, caspase-8-dependent and mitochondrion-independent pathway and the presence of an intact RIP homotypic interaction motif. Thereby, we suggested that TRIF-IRF3-IRF7 pathway activation, TRIF-NFκB pathway activation and apoptosis pathways are uncoupled and this can explain the different functions of TLR3-TRIF in neovascularization. Remarkably, we showed a regulation of pro-inflammatory cytokines via IRF3 and IRF7, which indicates that there may be a linkage between the TRIF-IRF and TRIF-NFκB pathways.
After phosphorylation downstream TLR3, IRF3 and IRF7 can produce type-I IFNs, such as IFNβ. However, the role of type-I IFNs and IRF3 and IRF7 in neovascularization is not identical. Schirmer et al observed attenuated post-ischaemic blood flow recovery 7 days after HLI in mice treated with type-I IFN, IFNβ. In addition, blocking IFNβ demonstrated to stimulate arteriogenesis via VSMC proliferation and IFnAR1−/− mice, deficient in the IFNβ receptor, improved blood flow recovery 7 days after induction of hind limb ischaemia. Teunissen et al observed that TRIF-NFκB-mediated pro-inflammatory cytokines are important regulators of neovascularization. These pre-clinical results are seemingly in contrast with our results since we show an impaired blood flow recovery after HLI in Irf3−/− and Irf7−/− mice. However, IRF3 and IRF7 do not only produce type-I IFNs, such as IFNβ, but can also regulate pro-inflammatory cytokine production, as we show in the gastrocnemius muscles of Irf3−/− and Irf7−/− mice. As we previously showed, TRIF-NFκB-mediated pro-inflammatory cytokines are important regulators of neovascularization.

The findings of the present study indicate that deficiency of IRF3 and IRF7 results in impaired post-ischaemic blood flow recovery caused by an attenuated angiogenesis and arteriogenesis linked to lack of inflammatory components in ischaemic tissue. In conclusion, IRF3 and IRF7 are essential regulators of neovascularization.

CONFLICT OF INTEREST STATEMENT

The authors confirm that there are no conflicts of interest.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

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