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Goel, Alexander N Gowda, Bhavani S Veena, Mysore S <u>et al.</u>

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Adipose-Derived Mesenchymal Stromal Cells Persist in Tissue-Engineered Vocal Fold Replacement in Rabbits

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	Objectives Cell therapies using mesenchymal stromal cells (MSCs) have been proposed as a promising new tool for the treatment of vocal fold scarring. However, the mechanisms by which MSC promote healing as well as their duration of survival within the host vocal fold have yet to be defined. The aim of this work was to assess the persistence of embedded MSCs within tissue-engineered vocal fold mucosal replacement in a rabbit model of vocal fold injury.
Abstract:	Methods Male rabbit adipose-derived mesenchymal stromal cells (ASC) were embedded within a three-dimensional fibrin gel, forming the cell-based outer vocal fold replacement (COVR). Four female rabbits underwent unilateral resection of vocal fold epithelium and lamina propria and reconstruction with COVR implantation. Polymerase chain reaction and fluorescent in situ hybridization for the sex-determining region of the Y chromosome (SRY-II) in the sex-mismatched donor-recipient pairs sought persistent cells after 4 weeks.
	Results A subset of implanted male cells was detected in the implant site at 4 weeks. Many SRY-II negative cells were also detected at the implant site, presumably representing native female cells that migrated to the area. No SRY-II signal was detected in contralateral control vocal folds.
	Conclusions The emergent tissue after implantation of a tissue-engineered outer vocal fold replacement is derived both from initially embedded adipose-derived stromal cells and infiltrating native cells. Our results suggest this tissue- engineering approach can provide a well-integrated tissue graft with prolonged cell activity for repair of severe vocal fold scars.

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9	6	AUTHORS:
10	7	Alexander N. Goel, BA ^a
11	8	Bhavani S. Gowda, PhD ^a
12	9	Mysore S. Veena, PhD ^b
13	10	Travis L. Shiba, MD ^a
14	11	Jennifer L. Long, MD, PhD ^{a,b} *
15		Jenniter L. Long, MD, FID
16	12	
17	13	^a Department of Head and Neck Surgery, David Geffen School of Medicine at University of
18 10	14	California, Los Angeles, Los Angeles, CA
19 20	15	^b Research Service, Greater Los Angeles Veterans Affairs Hospital System, Los Angeles, CA
20	16	
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37 38	31	*CORRESPONDING AUTHOR:
39 40	32	Dr. Jennifer L. Long
40	33	Email: <u>Jennifer.Long2@va.gov</u>
42	34	
43	35	David Geffen School of Medicine at UCLA
44	36	10833 Le Conte Ave
45	37	Los Angeles, Ca 90095
46	38	Ph: 310-825-4949
47	39	Fx: 310-206-7384
48	40	
49	41	Research Service
50	42	Greater Los Angeles VAHS
51		11301 Wilshire Blvd
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ABSTRACT

Objectives

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49	Cell therapies using mesenchymal stromal cells (MSCs) have been proposed as a promising new
50	tool for the treatment of vocal fold scarring. However, the mechanisms by which MSC promote
51	healing as well as their duration of survival within the host vocal fold have yet to be defined. The
52	aim of this work was to assess the persistence of embedded MSCs within a tissue-engineered
53	vocal fold mucosal replacement in a rabbit model of vocal fold injury.
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55	Methods
56	Male rabbit adipose-derived mesenchymal stromal cells (ASC) were embedded within a three-
57	dimensional fibrin gel, forming the cell-based outer vocal fold replacement (COVR). Four
58	female rabbits underwent unilateral resection of vocal fold epithelium and lamina propria and
59	reconstruction with COVR implantation. Polymerase chain reaction and fluorescent in situ
60	hybridization for the sex-determining region of the Y chromosome (SRY-II) in the sex-
61	mismatched donor-recipient pairs sought persistent cells after 4 weeks.
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63	Results
64	A subset of implanted male cells was detected in the implant site at 4 weeks. Many SRY-II
65	negative cells were also detected at the implant site, presumably representing native female cells
66	that migrated to the area. No SRY-II signal was detected in contralateral control vocal folds.
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68	Conclusions

- 69 The emergent tissue after implantation of a tissue-engineered outer vocal fold replacement is
- 70 derived both from initially embedded adipose-derived stromal cells and infiltrating native cells.
- 71 Our results suggest this tissue-engineering approach can provide a well-integrated tissue graft
- 72 with prolonged cell activity for repair of severe vocal fold scars.

KEYWORDS: Adipose-derived stromal cell; tissue engineering; vocal fold scarring; vocal fold

75 replacement

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INTRODUCTION

Vocal fold scarring, which can be caused by inflammation, radiation, trauma, or surgery, is a disabling voice disorder. Scar tissue alters the extracellular matrix (ECM) of the vocal fold, making it stiff and inelastic, leading to altered voice quality. The composition and organization of the ECM establish the vocal fold's biomechanical properties which in turn determine voice quality. Thus, treatment of the voice disturbance requires restoration of the unique microstructure of the ECM.

Cell-based approaches have received recent attention as potential therapies for vocal fold
scarring. Among various cell types, mesenchymal stromal cells (MSCs) are particularly
attractive given their ease of expansion in culture and extensive safety profile in humans.^{1,2}
MSCs can be administered alone or combined with a scaffold. Scaffolds provide an instructive
microenvironment that may enhance stem cell survival at the site of injury. We previously
showed that adipose-derived mesenchymal stromal cells (ASCs) embedded within a threedimensional fibrin gel scaffold improve vocal fold histology and vibration in rabbits.^{3,4}

Although MSC have been proposed for vocal fold scarring, the therapeutic mechanisms by
which MSC promote healing have yet to be defined. Reasons for benefit may include direct
replacement of damaged resident cells, release of immunomodulatory or trophic factors, or a
combination thereof. It is known that MSC secrete several anti-fibrotic and prosurvival factors
that likely play a role,⁵⁻⁷ but some evidence for differentiation and engraftment has also been
reported.⁸ The identification of MSC's powerful paracrine actions in other tissues has deemphasized long-term cell engraftment as a therapeutic requirement.⁹⁻¹¹ Nonetheless, implanted

cells must survive for some unknown duration to produce any paracrine effect. For vocal fold replacement, in which organized ECM remodeling is the therapeutic action, implanted cell influence should continue until remodeling is stabilized. Understanding the fate of transplanted stem cells requires reliable techniques for identifying cells after implantation. Traditional methods using stem cells labeled with transgenic constructs (e.g. green fluorescent protein) or membrane dyes (e.g. CM-DiI) have been used in the vocal fold with widely varying findings regarding the timeline of cell survival.¹²⁻¹⁴ These methods are limited by loss of gene expression in unstable transfections and reduction in the concentration of membrane dye through cell divisions.^{15,16} An alternative approach implants sex-mismatched ASCs and seeks signal for Y chromosome DNA.¹⁷ Such an approach capitalizing on donor-specific antigens may overcome some of the limitations of cell labeling.¹⁷ The aim of the present study is to assess the engraftment of ASCs within the female rabbit vocal fold implants that were previously reported. [3,4] Cell-based Outer Vocal fold Replacements (COVR) were formed with male rabbit ASC in three-dimensional fibrin gel scaffolds. COVRs were implanted in four female rabbits after removing the native vocal fold epithelium and lamina propria. Excised larynx phonation and histology results at 4 weeks were previously described. This study uses Y chromosome DNA to identify persistent donor cells four weeks after implantation, and TUNEL labeling to identify apoptotic cells. **METHODS** In Vitro Development of Cell-Based Outer Vocal Fold Replacement

Rabbit adipose-derived multipotent stromal cells (rASCs) were isolated from male rabbits and tissue-engineered constructs were created as previously described (Figure 1A).^{1,18,19} Briefly. rASCs were harvested from inguinal fat by collagenase digestion and expanded in culture. Cell multipotency was confirmed by differentiation to osteogenic and adipogenic phenotypes.¹⁸ Rabbit fibringen was mixed with bovine thrombin and rASC cell suspension to form fibrin gels with embedded ASC within 12mm Transwell culture inserts. The tissue constructs were cultured with an air interface and supplied culture medium through the insert base. Medium contained 10% fetal bovine serum and 100 ng/mL of epidermal growth factor. After 2 weeks, the cell-based outer vocal fold replacements (COVR) were harvested for implantation or baseline histology. Histologic stains included standard hematoxylin and eosin (H&E), Masson's trichrome to stain collagen fibers blue, and phosphotungstic acid hematoxylin (PTAH) to stain fibrin fibers blue. The mature COVR in vitro demonstrated homogeneous fibrin fibers, without PCL. detectable collagen. **COVR Implant Surgery** The local Institutional Animal Care and Use Committee approved this study which was conducted in accordance with all local and federal guidelines. Four female New Zealand white rabbits, weighing 3 to 3.5 kg, underwent survival surgery for outer vocal fold mucosa removal and implantation with COVR which was described previously.³ Briefly, rabbits were anesthetized, and the larynx exposed through a neck incision. After laryngofissure, the entire membranous cover layer was resected from the left inferior vocal fold. Immediately following

 resection, a mature COVR containing male rASC was placed to fill the defect and secured with

sutures and fibrin glue. All right vocal folds remained as untreated controls. The laryngofissure,

tracheotomy, and neck were closed and the animals recovered. Intramuscular dexamethasone,

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147	analgesics, and antibiotics were administered for three days to prevent laryngeal edema, pain,
148	and infection. All four animals recovered without complication or apparent distress, maintained
149	oral diet, and gained weight. After 4 weeks, animals were euthanized and larynges harvested.
150	The 4-week time point was chosen for best comparison with previous work in this laboratory and
151	because it lies within the range of other studies assessing MSC persistence. ^{3,4,8,13,20} All four
152	larynges produced phonation in an excised larynx setup, the same day as harvest as described
153	previously. ³ Larynges were then formalin-fixed and paraffin-embedded (FFPE) for sectioning
154	and DNA extraction.

156 PCR Amplification of Rabbit Y Chromosome DNA

157 Genomic DNA from male New Zealand white rabbit thymuses was isolated and purified using 158 the Qiagen genomic DNA kit (Germantown, MD). Primers were constructed for the second sexdetermining region of the Y chromosome (SRY-II) of the European brown hare.²¹ The isolated 159 160 DNA was amplified by PCR using the SRY-II primers with AmpliTag Gold[™] 360 Master Mix 161 (Applied Biosystems, Foster City, CA) under the following conditions: 95°C for 2 minutes 162 followed by 35 cycles of 95°C for 45 seconds, 55°C for 30 seconds, 72°C for 45 seconds with a 163 final extension at 72°C for 5 minutes. PCR products were run on 1.5% agarose gel and images 164 were captured using G:BOX chemi XRQ (Syngene, Frederick, MD) to confirm expected SRY-II 165 size of 670 base pairs.

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After validating the primer sequence with male rabbit DNA, the primers were used to amplifyDNA from rabbit larynx tissues. Implanted female rabbit vocal fold tissues were dissected out of

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3 4	169	the paraffin blocks, using a scalpel to sharply divide the implanted (left) and control (right) vocal
5 6	170	folds at the anterior commissure (as indicated in Figure 1B). DNA was isolated and purified
7 8 9	171	using the QIAamp DNA FFPE tissue kit (Qiagen, Germantown, MD). PCR amplification
9 10 11	172	proceeded as above using SRY-II primers. PCR products were separated on 1.5% agarose gel
12 13	173	and imaged.
14 15	174	
16 17 18	175	Fluorescent In Situ Hybridization (FISH)
19 20	176 177	DNA was isolated from normal male rabbit thymus, and Y chromosome-specific DNA was then
21 22	178	amplified as described above. PCR products were separated on 0.8% agarose gel. The amplified
23 24 25 26 27 28 29	179	670 bp SRY-II product was excised from the gel and purified using QIAquick Gel Extraction kit
	180	(Qiagen, Germantown, MD). Purified DNA was labeled with Alexa Fluor 647 dye using Fish
	181	Tag DNA multicolor kit (Molecular Probes, Eugene, OR) following the protocol provided with
30 31	182	the kit.
32 33	183	
34 35 36	184	Five-micron FFPE tissue sections of implanted rabbit larynges were used for FISH. Y
30 37 38	185	chromosome-specific rabbit DNA labeled with Alexa Fluor 647 was used as FISH probe. Tissue
39 40	186	sections fixed on glass slides were de-paraffinized in Xylene (with three changes of 5 minutes
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43	187	each), then gradually rehydrated using 100%, 85%, and 50% ethanol (one minute each) followed
44 45 46	188	by incubation in 0.2 N HCl for 20 minutes, distilled water for 10 minutes, 2X sodium-saline
46 47 48	189	citrate for two minutes, 10mM sodium citrate buffer pH 6.0 at 80°C for 30 minutes, 20 ng
49 50	190	proteinase K/ml in 10 mM HCl at 37°C for 20 minutes and finally in distilled water for three
51 52	191	minutes. Slides were then dehydrated gradually in 50%, 70%, 85% and 100% ethanol (one
53 54	192	minute each) and air dried for 30 minutes. For the rest of the procedures for pre-hybridization,
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multicolor kit user guide (Invitrogen, Carlsbad, CA). Slides were coverslipped and imaged with a

hybridization, and post-hybridization, washes were carried out as described in the Fish Tag

647 nm fluorescence microscopy filter.

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97	Apoptosis Assay
98	Terminal deoxynucleotidyl transferase dUTP nick-end labeling (TUNEL) assay for apoptotic
99	cells was performed on paraffin-embedded, formalin-fixed slides from implanted rabbit larynges
00	(Trevigen TACS 2TdT DAB labeling kit, Gaitersburg, MD). The terminal deoxynucleotidyl
01	transferase (TdT) enzyme incorporated biotin label into fragmented DNA of apoptotic cells.
02	Positive controls were treated with TACS-Nuclease enzyme to induce DNA strand nicks. Biotin
03	was detected with diaminobenzidine (DAB) substrate, and nuclei counterstained with methyl
04	green. Images were taken at 40x.
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06	RESULTS
07	Rabbit Y chromosome PCR
08	Thymic DNA from normal male New Zealand white rabbits confirmed amplification with the
09	SRY-II primer sequences, producing a 670 bp PCR product.
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11	DNA was isolated from implanted and control rabbit vocal folds, amplified with SRY-II primers,
12	and separated on agarose. All four implanted vocal folds (with male cells applied to female
13	larynx) showed an amplified DNA band that corresponded with the 670 bp PCR product from
14	male thymus. All four contralateral control vocal folds (female larynx without implant) lacked
15	any amplification with the SRY-II primer (Figure 2A).

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5 6	217	Fluorescent In Situ Hybridization (FISH)
7 8 9 10 11	218	In situ hybridization with a tagged SRY-II probe confirmed the findings of PCR. In all four
	219	female larynges, scattered positive (male) cells were detected on the operated side. The area of
12 13	220	implantation also contained many cells negative for the Y-chromosome signal that presumably
14 15 16	221	represented native female cells that had migrated into the implanted site (Figure 2B). Implanted
16 17 18	222	ASCs were primarily localized to the lamina propria, with relatively fewer in the epithelial layer
19 20	223	and occasional pockets detected in deep tissue near the thyroarytenoid muscle. No SRY-positive
21 22	224	cells were detected in contralateral control vocal folds.
23 24 25	225	
26 27 28 29 30 31 32 33 34 35 36 37 38 39	226	Histological Examination
	227	Figure 1B shows H&E staining, with implanted left vocal fold appearing similar to unoperated
	228	right vocal fold. Collagen fibers were previously demonstrated in the region of implantation on
	229	Masson's trichrome stain, indicating remodeling of the provisional scaffold. ³ Phosphotungstic
	230	acid hematoxylin (PTAH) staining was attempted to determine persistence of the fibrin matrix,
	231	but the non-specific nature of the staining was judged to be inconclusive.
39 40 41	232	
41 42 43 44 45 46 47 48 49 50	233	Apoptosis assay
	234	TUNEL labeling for apoptotic cells was negative within the area of COVR implant, consistent
	235	with its negative labeling in contralateral unoperated vocal folds. Positive controls of nuclease
	236	enzyme-treated implant slides did demonstrate positive TUNEL staining, confirming the labeling
51 52	237	technique (Figure 3).
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DISCUSSION

In this study, we investigated the engraftment of rabbit ASCs within a tissue-engineered implant in the rabbit vocal fold. To distinguish transplanted cells from native cells, male rASCs were seeded into female hosts to permit DNA analysis for genes on the Y-chromosome. Four weeks after implantation, scattered male cells were localized within the implant site, along with many cells that were negative for the fluorescence hybridization signal. This finding suggests that the emergent tissue derives both from infiltrating native cells and implanted ASCs that were initially seeded onto the scaffold.

Simple injections of mesenchymal stromal cells have prevented scar formation in previous animal models.^{20,22,23} Postulated mechanisms include secretion of growth factors active in vocal fold tissue repair, including hepatocyte growth factor (HGF) and fibroblast growth factor 2 (FGF2).^{7,24,25} In addition, Hiwatashi et al reported that MSC treatment led to increased host expression of hyaluronan synthase, which is thought to maintain viscoelastic function of the vocal folds.¹² Transient cell activity may be adequate to prevent scarring when the cells are applied concordant with injury. However, accumulating data have revealed that the survival rate of cells injected into previously scarred vocal folds is disappointingly low. Svensson et al did not detect any GFP-positive MSC three months after direct administration in scarred rabbit vocal folds.²⁶ de Bonnecaze studied an earlier 3-week time point and also were unable to detect GFP-labeled cells after MSC injection.¹³ Survival of transplanted cells is limited by apoptosis, lack of trophic factors, pathologic conditions, and the inflammatory response.

Page 13 of 21

Scaffolds, including hyaluronic acid (HA) based hydrogels^{27,28} and atelocollagen,²² have been used as delivery carriers and led to increased cell survival in a few prior studies. These scaffolds can be engineered to mimic the native extracellular matrix which may create a microenvironment more suitable to MSC survival. HA scaffolds have been most widely employed, but are limited by challenges in preventing HA degradation and clearance in vivo that allows cells to migrate rapidly away from the site of implantation.²⁹ In the largest pre-clinical trial for vocal fold scar, Bartlett et al investigated cell engraftment after administration of MSC with and without a hyaluronic acid gel carrier in rabbits and detected no transplanted cells in either group at 2 weeks.³⁰ In contrast, our study utilized fibrin as the scaffold matrix and detected implanted cells at 4 weeks in all four female rabbits. Protein-based scaffolds, such as fibrin, may be superior in promoting cell-to-surface binding compared to polysaccharide-based scaffolds such as HA, potentially enhancing cell survival.³¹ We hypothesize that the fibrin scaffold provides a supportive niche which improves cell survival relative to cell injection alone. Whether the surviving ASCs establish a long-term, stable sub-population residing within the scaffold cannot be adequately determined by our 4-week study, but this is a question worthy of further investigation. In this study, host cells did significantly infiltrate the implant, which by 4 weeks was more host-derived than graft-derived.

279 Characterizing the fate of transplanted cells in regenerative medicine applications depends on 280 having reliable techniques for cell tracking or detection. To date, methods to identify implanted 281 stem cells in the vocal fold have primarily relied on transgene construct labeling (e.g. green 282 fluorescent protein) or membrane dye approaches. These methods are attractive given their ease 283 of use, but suffer potential limitations in the long-term evaluation of stem cell persistence. The

GFP transgene may become silenced as cells proliferate and differentiate *in vivo*, potentially masking cell detection.^{15,16,30} Alternatively, lipophilic membrane labels such as carbocyanine derivatives may be detected even after macrophage ingestion of dead cells, potentially leading to overestimation of surviving cell populations.³² Here, we target sex-specific antigens existing exclusively in the injected cells by immunofluorescence in a sex-mismatched animal model. Such a technique may overcome the limitations of cell labeling methods without a significant increase in complexity or cost and may be a desirable approach for future studies. The limitation of this approach is that it only applies to male cells in female hosts. Because this methodology of Y-chromosome detection is invalid in the male-male transplant, the cell persistence in male hosts remains unknown. Inflammatory cell infiltrate did occur in the female hosts but not in male hosts receiving allogeneic male cells.³ This study transplanted allogeneic ASCs in the absence of immunosuppression, and the persistence of some implanted cells after 4 weeks indicates the inflammatory response did not reach the level of complete transplant rejection. ASCs themselves have significant immunomodulatory activity, which could assist implanted cells in evading the host immune system.^{33,34} Why the females exhibited greater leukocytosis is of interest for further investigation. It is conceivable that the female host has inherently greater immune activity than the males. Or, the greater female response may simply be due to greater antigen mismatch with male cells. In studies of solid organ transplant in humans, it has been observed that sex mismatch in donor-recipient pairs, especially cases of male donor to female recipient, does lead to higher incidence of acute rejection.^{35,36} The mechanism has been linked to the production of alloantibodies against the male H-Y minor histocompatibility complex antigen.³⁷ Very limited basic work in cell therapies has explored sex

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3 4	307	as a biological variable, which is essential to distinguish these hypotheses in future research. ³⁸
5 6	308	For eventual clinical applications, the use of autologous or bank-matched stem cells would be
7 8 9	309	expected to further reduce immune response which could prolong cell survival.
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12 13	311	CONCLUSION
14 15 16	312	This study furthers the pre-clinical development of a tissue-engineered outer vocal fold
17 18	313	replacement constructed from adipose-derived stromal cells. A subset of the implanted cells
19 20	314	persisted at least 4 weeks in rabbits, in contrast to other studies with different implant designs.
21 22 23	315	Our results suggest this tissue-engineering approach can provide a well-integrated tissue graft
23 24 25	316	with prolonged cell activity for repair of severe vocal fold scars.
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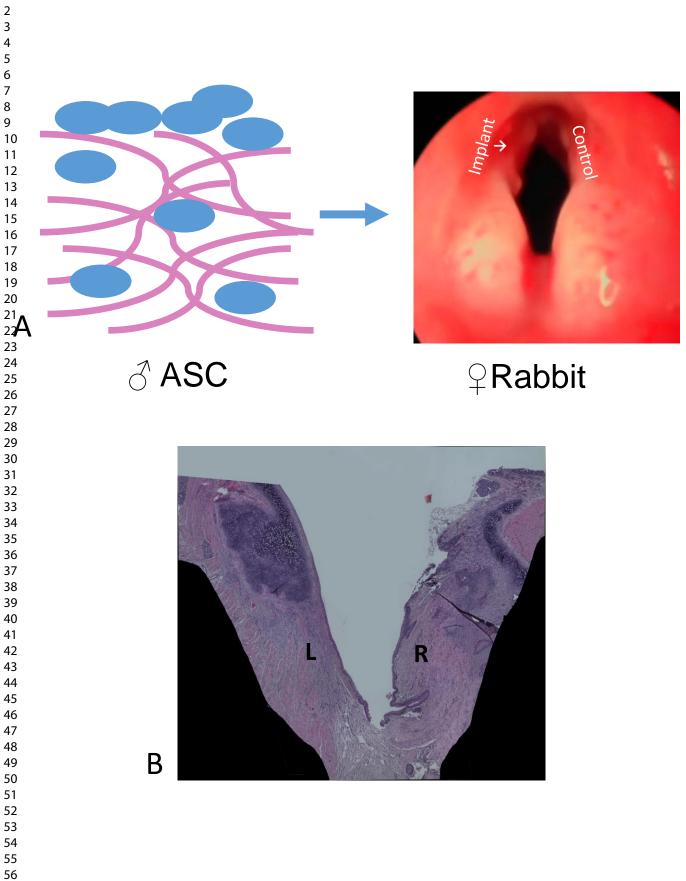
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2 3 4	422	FIGURE LEGENDS
5 6 7 8 9	423	Figure 1. (A) Donor male rASCs were seeded in a COVR that was surgically implanted to
	424	female rabbit recipients. (B) Hematoxylin and eosin (H & E) stain of harvested larynx, where left
9 10 11	425	vocal fold received COVR implant (L) and right vocal fold served as unoperated control (R).
12 13	426	10x. (Rabbit 4). Thick black line at the anterior commissure indicates the dividing line for
14 15	427	extracting DNA from individual vocal folds.
16 17 18	428	
18 19 20	429	Figure 2. Presence of implanted cell DNA after 1 month. (A) PCR amplification of SRY-II
21 22 23 24	430	DNA sequence. "SRY control": normal male rabbit tissue. Lanes are shown for each rabbit's
	431	implanted vocal fold ("Implant") and contralateral non-implanted vocal fold ("Contra"). (B)
25 26 27	432	Fluorescent in-situ hybridization for Y chromosome DNA (SRY-II probe, appearing red) in
28 29	433	female rabbit vocal folds 4 weeks after implantation of a COVR seeded with male ASCs. Signal
30 31	434	appears in the operated left vocal folds (L), but not in the contralateral unoperated right vocal
32 33 34	435	folds (R), suggesting engraftment of implanted ASCs. Nuclei are stained with DAPI (blue).
35 36	436	Rabbit 1 at 4X magnification, Rabbit 2 at 10X magnification.
37 38	437	
39 40	438	Figure 3. TUNEL labeling for apoptotic cells. (A) Positive control for TUNEL staining,
41 42 43	439	enzymatically treated to induce DNA nicks. R4, 40x (B) Harvested left vocal fold (COVR
44 45	440	implant side) with negative TUNEL staining confirmed that the cells were not undergoing
46 47	441	apoptosis. Rabbit 3, 40x. (C) Harvested right vocal fold (unoperated control) also with negative
48 49 50	442	TUNEL staining, as expected. Rabbit 3, 40x.
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Page 21 of 21

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