



Article

Atrial Fibrillation in Heart Failure Is Associated With High Levels of Circulating microRNA-199a-5p and 22–5p and a Defective Regulation of Intracellular Calcium and Cell-to-Cell Communication

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SUPPLEMENTARY TABLES

Table S1. Characteristics of the first cohort.

| | permAF (<i>n</i> = 9) | SR (<i>n</i> = 9) | <i>p</i> -Value |
|--------------------------------|------------------------|--------------------|-----------------|
| Patient Characteristics | | | |
| Sex, men (%) | 5 (55.5) | 5 (55.5) | NS |
| Age, y (SD) | 74.9 (8.2) | 75.2 (7.0) | NS |
| BMI, kg/m ² (SD) | 26.8 (2.6) | 26.8 (2.8) | NS |
| Medical history | | | |
| Tobacco history (%) | 4 (44.4) | 5 (55.5) | NS |
| Hypertension (%) | 7 (77.8) | 7 (77.8) | NS |
| Diabetes (%) | 2 (22.2) | 6 (66.7) | NS |
| Hypercholesterolemia (%) | 4 (44.4) | 7 (77.8) | NS |
| COPD (%) | 2 (22.2) | 1 (11.1) | NS |
| CKD (%) | 2 (22.2) | 2 (22.2) | NS |
| Previous stroke (%) | 2 (22.2) | 2 (22.2) | NS |
| HF parameters | | | |
| Ischemic etiology (%) | 4 (44.4) | 4 (44.4) | NS |
| Heart rate, bpm (SD) | 70.2 (13.4) | 80.0 (16.3) | NS |
| NYHA class (%): | | | NS |
| - I-II | 6 | 6 | |
| - III-IV | 3 | 3 | |
| LVEF, % (SD) | 32.0 (6.5) | 28.1 (5.6) | NS |
| Log Pro-BNP, mean (SD) | 3.4 (0.5) | 3.2 (0.6) | NS |
| HF treatment | | | |
| ACEI/ARB (%) | 8 (88.9) | 8 (88.9) | NS |
| Beta-blockers (%) | 9 (100) | 9 (100) | NS |
| Diuretics (%) | 9 (100) | 9 (100) | NS |
| Heart rhythm | | | |
| Permanent SR | 0 | 9 | |
| History of parox/pers AF* | 0 | 0 | |
| Permanent AF# | 9 | 0 | |

BMI: body mass index; COPD: chronic obstructive pulmonary disease; CKD: chronic kidney disease; NYHA: New York Heart Association; LVEF: left ventricular ejection fraction; ACEI: angiotensin-converting-enzyme inhibitors; ARB: angiotensin-receptor blockers; SR: sinus rhythm; AF: atrial fibrillation. *Includes episodes of paroxysmal or short-standing persistent AF; #includes long-standing (> 1 year) persistent or permanent AF. NS: non-significant.

Table S2. MicroRNAs differentially expressed in the discovery phase.

| miRNA | <i>p</i> -Value | miRNA | <i>p</i> -Value | miRNA | <i>p</i> -Value |
|--------------------|-----------------|-------------------|-----------------|-----------------|-----------------|
| miR-505# | 0.000701 | miR-1 | 0.010444 | miR-140 | 0.024151 |
| miR-20a | 0.002003 | miR-151-5P | 0.010454 | miR-671-3p | 0.024503 |
| miR-539 | 0.002037 | miR-645 | 0.011216 | miR-664 | 0.024804 |
| miR-425# | 0.002114 | miR-7# | 0.011294 | miR-195 | 0.025214 |
| miR-106a | 0.002469 | miR-93# | 0.011334 | miR-9# | 0.025847 |
| miR-30d# | 0.002633 | miR-130b# | 0.012461 | miR-124a | 0.026115 |
| miR-19a | 0.004109 | miR-186 | 0.012493 | miR-495 | 0.026444 |
| miR-16 | 0.004259 | miR-652 | 0.012506 | miR-151-3p | 0.026993 |
| miR-628-3p | 0.004991 | miR-26a | 0.012758 | miR-106b | 0.029477 |
| miR-199a-5p | 0.006002 | miR-432 | 0.014040 | miR-340 | 0.030723 |
| miR-301 | 0.006242 | let-7d | 0.014543 | miR-185 | 0.033889 |
| miR-374 | 0.006285 | miR-23a | 0.014962 | miR-411 | 0.035389 |
| miR-133a | 0.006676 | miR-636 | 0.015891 | miR-590-3P | 0.035525 |
| miR-374-5p | 0.007119 | miR-20a# | 0.016181 | miR-98 | 0.035920 |
| miR-330 | 0.007449 | miR-324-5p | 0.016251 | miR-30e-3p | 0.035936 |
| miR-126# | 0.007465 | miR-191# | 0.016514 | miR-1180 | 0.036382 |
| miR-17 | 0.007740 | miR-140-3p | 0.016646 | miR-27a | 0.037929 |
| miR-142-5p | 0.007847 | miR-625 | 0.017292 | miR-331-5p | 0.040697 |
| miR-1255B | 0.008360 | miR-483-3p | 0.017459 | miR-656 | 0.042341 |
| miR-598 | 0.008435 | miR-744 | 0.018172 | miR-107 | 0.043088 |
| miR-26a-1# | 0.008641 | miR-489 | 0.018316 | miR-331 | 0.043250 |
| miR-27b | 0.008667 | miR-24 | 0.019166 | miR-326 | 0.043429 |
| miR-22-5p | 0.008810 | miR-28 | 0.019784 | miR-146b | 0.043554 |
| miR-148b | 0.009361 | miR-380-3p | 0.021854 | miR-221 | 0.046365 |
| miR-191 | 0.010162 | miR-580 | 0.023425 | miR-15b | 0.048038 |
| miR-125a-5p | 0.010217 | | | | |

MiRNA selected for the replication phase are marked in bold.

Table S3. Characteristics of the donors of atrial tissue.

| Tissue Collection Site | LAA | | RAA | |
|---|-------------|-------------|-------------|-------------|
| | No AF | AF | No AF | AF |
| Total number of patients | 6 | 6 | 5 | 5 |
| Mean age, years (\pm SD) | 60 \pm 14 | 66 \pm 12 | 64 \pm 10 | 68 \pm 12 |
| Men, <i>n</i> (%) | 4 (67) | 5 (83) | 4 (80) | 5 (100) |
| Surgical procedure, <i>n</i> (%) | | | | |
| CABG \pm AVR/MVR | 2 (33) | 1 (17) | 2 (40) | 1 (20) |
| AVR/MVR | 4 (67) | 5 (83) | 3 (60) | 4 (80) |
| Medical history, <i>n</i> (%) | | | | |
| Smoker/ex-smoker | 4 (67) | 2 (33) | 3 (60) | 1 (20) |
| Hypertension | 3 (50) | 3 (50) | 2 (40) | 3 (60) |
| Diabetes mellitus | 2 (33) | 0 (0) | 0 (0) | 0 (0) |
| Previous MI | 1 (17) | 2 (33) | 0 (0) | 0 (0) |
| COPD/asthma | 0 (0) | 0 (0) | 0 (0) | 0 (0) |
| Medications, <i>n</i> (%) | | | | |
| Anticoagulants | 1 (17) | 4 (67) | 0 (0) | 3 (60) |
| β -Blockers | 2 (33) | 5 (83) | 3 (60) | 5 (100) |
| Statins | 2 (33) | 3 (50) | 1 (20) | 3 (60) |
| Calcium-channel blockers | 0 (0) | 0 (0) | 0 (0) | 1 (20) |
| ACEIs and ARBs | 1 (17) | 4 (67) | 1 (20) | 3 (60) |
| Diuretics | 0 (0) | 2 (33) | 2 (40) | 2 (40) |

LAA, left atrial appendage; RAA, right atrial appendage; noAF, sinus rhythm; AF, atrial fibrillation; ACEI, angiotensin-converting enzyme inhibitor; ARB, angiotensin II receptor blocker; AVR, aortic valve replacement; CABG, coronary artery bypass surgery; COPD, chronic obstructive pulmonary disease; MI, myocardial infarction; MVR, mitral valve replacement. Percentage in parentheses (%) indicates percentage within the same group.

Table S4. MiRNA primer sequences for atrial tissue.

| No AF | AF |
|-------|----|
|-------|----|

| Assay Name | Assay ID | Target Sequence 5'-3' |
|-----------------|------------|-------------------------|
| hsa-miR-22-5p | 477987_mir | AGUUCUUCAGUGGCAAGCUUUA |
| hsa-miR-199a-5p | 478231_mir | CCCAGUGUUCAGACUACCUGUUC |
| hsa-miR-191-5p | 477952_mir | CAACGGAAUCCAAAAGCAGCUG |
| hsa-miR-26a-5p | 477995_mir | UUCAAGUAUCCAGGAUAGGCU |

SUPPLEMENTARY FIGURES

FIG.S1

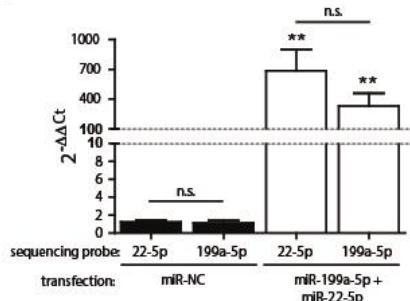


FIG.S2

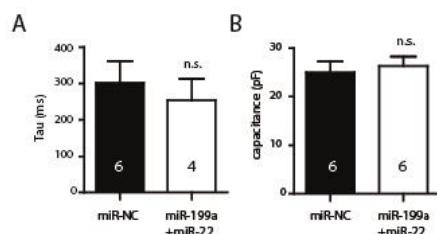


FIG.S3

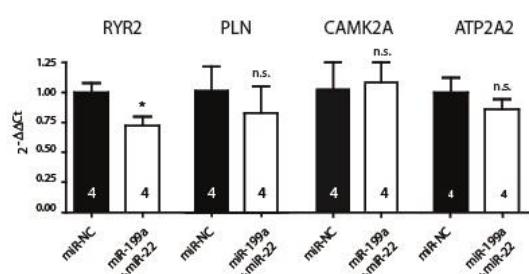


Figure S1. RNA expression levels of miR-22-5p and miR-199a-5p in HL1 cells transfected with miR-199a+miR-22 or miR-NC. Specific probes for each miRNA were used. Data are expressed as the fold change ($2^{-\Delta\Delta Ct}$) relative to the negative control, miR-NC. **Figure S2.** A. Fast inactivation kinetics (τ_{inact}) of I_{CaL} currents and, B. cell capacitance of HL-1 cells transfected with miR-199a+miR-22 (white) or miR-NC (black). Numbers within the columns represent number of patched cells. **Figure S3.** RYR2, PLN, CAMK2A and ATP2A2 mRNA expression levels in miR-199a+miR-22 HL-1 cells, normalized to miR-NC. n.s.: non-significant, * $p < 0.05$, ** $p < 0.005$.