Transplantation of autologous extracellular vesicles for cancer-specific targeting.

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Supplementary Materials



Figure S1. Representative EV imaging of PDEVs by cryo-electron microscopy, scale bar: 100nm.



Figure S2. Comparison of NTA profiles of patient-derived EVs before (blue line) and after (red line) ICG loading. The graph reports the mean (solid line) and the S.E.M. (dotted line) of n=6 measures for each experimental group. ICG encapsulation did not significantly change the size distribution profile of EV nanoparticles.



Figure S3. Pseudo-color representative images of ICG fluorescence. *In-vivo* and *ex vivo* imaging of MC-38 tumor-bearing mice (black mice) or mice bearing mammary cancer (NeuT, white mice) injected with EVs obtained from heathy volunteers (left panels) or patient-derived EVs (right panels), each labeled with ICG. Patients-derived EVs showed tropism for MC-38 tumor and for the mammary cancer originated in the NeuT mice: a heterologous (independent of the tumor type) recognition ability of neoplastic tissue as previously reported for cancer-derived EVs (see reference 16 in the main text). No signal could be detected in both mouse model of cancer when injected with ICG-labelled EVs isolated from healthy volunteers.



Figure S4. (A) LC chromatographic profiles of protein content in preparations of EVs from CRC patients (left panel) or healthy donors, did not show significant differences in the abundance of the most represented proteins. (B) MS quantification of albumin and lipoprotein content in CRC patients (blue bars) or healthy donors (red bars) did not detect significant differences between the two groups, as assessed by statistical analyses reported in the right panel.



Figure S5. Pseudo-color representation of ICG fluorescence. *In vivo* (left panels) and *ex vivo* (right panels) imaging of PDX mice injected with *i*) patient-derived EVs labeled with ICG (EV-ICG) or *ii*) free ICG ((130 nmol, the same amount included as cargo into the patient-derived EVs).



Figure S6. Picture of tumor tissue dissected from PDX II mouse, acquired by confocal laser scanning microscopy (CLSM) and showing the fluorescent signals of nuclei stained with DAPI (blue) and the fluorescence released by ICG (red). In order to assess the distribution of ICG fluorescence in the tissue sections, optical sectioning was performed by acquiring a series of focal planes distant 1 μ m along the optical (z) axis of the microscope, using the z-scan mode available on Nikon A1R laser scanning confocal microscope. Orthogonal projections were then reconstructed using ImageJ software (NIH). Briefly, the software projected on the y axis the fluorescent signal acquired in each optical section, simulating a lateral view of the tissue slice to reveal the interior configuration of the imaged region (right panel). (Scale bar: 100 μ m)



Figure S7. Low-power view of canine mammary-gland tumor specimen classified as carcinoma, represented as a superposition of bright-field (gray) and ICG fluorescence (red). The picture shows the accumulation of fluorescent signal in the tumor tissue (T). Details of the tumor area are reported in Fig. 6B. Scale bar: 2 mm.



Figure S8. Superposition of bright-field (gray) and ICG fluorescence (red) microscopy images, captured by confocal microscopy from canine specimens obtained from a tumor (mastocytoma) patient. Picture represents the resection margin within a region of normal tissue (A) or region within the mastocytoma tissue (B) of the same patient. S = skin; SC = subcutis; T = tumor (Mast Cell Tumor). Scale bar: 2mm. While samples collected in the non-tumor, cancer-free tissue did not show detectable specific fluorescent emission (A), neoplastic tissue (T) exhibited fluorescence in the NIR spectrum related to ICG (B) thus corroborating the evidence that, also in canine large breeds with spontaneous tumors, a specific labeling of the autologous neoplastic tissue can be obtained with the fluorescent dye conveyed by PDEVs.

 Table S1. Patients and tumors characteristics of CRC-bearing patients recruited for PDX generation.

| Patient number | Age | Sex | Primary tumor (previously removed) | Stage | Histology Genotype | Serum Markers |
|-------------------|-----|-----|---|---|---|---------------------------|
| I | 51 | F | Left colon (sigmoid) | pT4a, N1a, M1 Synchronous liver M+ | Adenocarcinoma G2 RAS/BRAF wild type | CEA: 1.5 CA 19.9: 24 |
| п | 63 | М | Right colon | pT1, N1, M0 Metachronous liver M+ | Adenocarcinoma G2 KRAS: mutated (G12V) | CEA: 5.69 |
| ш | 69 | М | Right colon | pT3, N1a, M1 Synchronous liver M+ | Adenocarcinoma G2 NT3, N1a, M1 Nronous liver M+ (Codon 13) | |
| IV | 73 | F | Right colon | pT4, N0, M0 Metachronous liver M+ | Adenocarcinoma G2 KRAS wild type | CEA: 1.5 |
| v | 54 | F | Left colon (sigmoid) | pT2, N0, M0 Metachronous liver M+ | Adenocarcinoma G2-G3 KRAS wild type | CEA: 2.05 CA 19.9: 8.0 |
| VI | 70 | F | Right colon | pT3, N0, M1a Synchronous liver M+ | Adenocarcinoma G2 RAS/BRAF wild type | CEA:2.84 CA 19.9: 18.4 |
| VII | 77 | F | Right colon | pT3, N1, M1 Synchronous liver M+ | Adenocarcinoma G3 BRAF mutated (V600E) KRAS/ NRAS wild type | CEA: 5.2 |
| VIII | 70 | М | Left colon (sigmoid) | pT3, N2, M1 Synchronous liver M+ | Adenocarcinoma G2-G3 RAS/BRAF wild type | CEA: 51 |

Table S2. Results of blood tests on specific panels for testing bone marrow, liver and kidney functions before and after EV injection. No significant variations were observed at early intervals after ICG-loaded EVs.

| | | Pre- IC EVs i | CG-loaded njection | Post- ICG-loaded EVs injection (24hrs) | |
|------------|-------|------------------|-----------------------|---|-------|
| Test | Unit | Dog 1 | Dog 2 | Dog 1 | Dog 2 |
| RBC | n/µL | 6.23 | 5.06 | 6.69 | 4.53 |
| WBC | n/µL | 7.75 | 16.41 | 7.70 | 8.83 |
| ALP | U/L | 74 | 231 | 71 | 197 |
| ALT | U/L | 57 | 70 | 48 | 67 |
| Creatinine | mg/dL | 1.3 | 0.5 | 1.0 | 0.4 |
| Urea | mg/dL | 64 | 17 | 36 | 24 |