

1 **Sennoside A, derived from the Traditional Chinese Medicine plant**  
2 ***Rheum L.*, is a new dual HIV-1 inhibitor effective on HIV-1 replication**

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### 34 **ABSTRACT**

35 **Background:** Despite the availability of effective antiretroviral therapies, drugs  
36 for HIV-1 treatment with new mode of action are still needed. An innovative  
37 approach is aimed to identify dual HIV-1 inhibitors, small molecules that can  
38 inhibit two viral functions at the same time. Rhubarb, originated from *Rheum*  
39 *palmatum* and *R. officinale*, is one of the earliest and most commonly used  
40 medicinal plants in Traditional Chinese Medicine (TCM) practice. We wanted  
41 to explore TCM for the identification of new chemical scaffolds with dual action  
42 abilities against HIV-1.

43 **Methods:** *R. palmatum* and *R. officinale* extracts along with their main single  
44 isolated constituents anthraquinone derivatives were tested on both HIV-1  
45 Reverse Transcriptase (RT)-associated DNA Polymerase (RDDP) and  
46 Ribonuclease H (RNase H) activities in biochemical assays. Active  
47 compounds were then assayed for their effects on HIV-1 mutated RTs,  
48 integrase (IN) and viral replication.

49 **Results:** Both *R. palmatum* and *R. officinale* extracts inhibited the HIV-1  
50 RT-associated RNase H activity. Among the isolated constituents, Sennoside  
51 A and B were effective on both RDDP and RNase H RT-associated functions in  
52 biochemical assays. Sennoside A was less potent when tested on K103N,  
53 Y181C, Y188L, N474A and Q475A mutated RTs, suggesting the involvement  
54 of two RT binding sites for its antiviral activity. Sennoside A affected also HIV-1

55 IN activity *in vitro* and HIV-1 replication in cell-based assays. Viral DNA  
56 production and time of addition studies showed that Sennoside A targets the  
57 HIV-1 reverse transcription process.

58 **Conclusion:** Sennoside A is a new scaffold for the development of HIV-1 dual  
59 RT inhibitors.

60

61 **Keywords:** Sennoside A, HIV-1, Reverse Transcriptase, antiviral activity, plant  
62 extracts, dual inhibitor.

63

64 **Abbreviations:** DKA, diketo acid; EC<sub>50</sub>, HIV-1 replication by 50%; HAART,  
65 Highly Active Anti-Retroviral Therapy; HIV-1, Human Immunodeficiency Virus  
66 type 1; IN, integrase; INIs, HIV-1 IN inhibitors; NNRTI, Non-Nucleoside RT  
67 Inhibitor; NRTIs, Nucleoside RT Inhibitors; PI, Protease Inhibitor; qPCR,  
68 quantitative real-time PCR; RDDP, Reverse Transcriptase (RT)-associated  
69 DNA Polymerase; RNase H, Ribonuclease H; TCID<sub>50</sub>50%/ml, Tissue Culture  
70 Infective Doses; TCM, Traditional Chinese Medicine.

71

## 72 1. INTRODUCTION

73 The Human Immunodeficiency Virus type 1 (HIV-1) encoded Reverse  
74 Transcriptase (RT) is responsible for the reverse transcription process of the  
75 viral genome, a fundamental step in the HIV-1 replication cycle and a still  
76 attractive target for the identification of new antiretroviral inhibitors (Esposito et  
77 al. 2012; Corona et al. 2013; Distinto et al. 2013; Esposito and Tramontano  
78 2013). The HIV-1 RT is a multifunctional protein endowed of two main  
79 enzymatic functions: an RNA-Dependent DNA Polymerase (RDDP) activity,  
80 which accounts for the formation of the RNA:DNA intermediate, and a  
81 Ribonuclease H (RNase H) activity, involved in the hydrolytic cleavage of the  
82 RNA strand of the RNA:DNA hybrid (Esposito et al. 2012). Currently, for the >  
83 30 million of people worldwide infected by HIV-1 no vaccine is available, but an  
84 effective antiretroviral therapy has been developed, termed Highly Active  
85 Anti-Retroviral Therapy (HAART). HAART first line treatment is usually  
86 composed by a combination of two Nucleoside RT Inhibitors (NRTIs) to which  
87 a Non-Nucleoside RT Inhibitor (NNRTI) or a Protease Inhibitor (PI) is added.  
88 HIV-1 IN inhibitors (INIs) are expected to retain efficacy against HIV-1 strains  
89 resistant to other antiretroviral drugs, and represent a valid second-line  
90 therapy. The acute and chronic drug toxicities and the selection of drug  
91 resistant strains, in particular to RTIs, represent a valid reason to identify new  
92 antiviral agents. Indeed, the development of drugs with new mode of action,  
93 such as single molecules that may act on two different target enzymes or two

94 different catalytic functions, would reduce the number of administered drugs,  
95 their chronic toxicity and the chance of selecting drug resistant viruses  
96 ( Distinto et al. 2013; Esposito and Tramontano 2013) .

97 In the last 20 years, several research efforts have been focused on exploiting  
98 natural products as scaffolds for the development of potential antiviral agents  
99 (Li and Vederas 2009; Cos et al. 2008; Yu et al. 2007; Bicchi et al. 2009; Xu et  
100 al. 2015), although no natural compound able to block HIV-1 replication has  
101 ever entered clinical trials. In China, Traditional Chinese Medicine (TCM) is the  
102 major ancient therapeutic system and its herbal component is the most  
103 important one. Radix et Rhizoma Rhei (Da Huang in Chinese), usually  
104 originated from *Rheum palmatum* and *R. officinale*, is one of the earliest and  
105 also the most commonly used medicinal plant in TCM practice (Beijing:  
106 Chinese Pharmaceutical Science and Technology Publishing House, 2010.

107 The earliest record of Rhubarb use appears to come from the Shen Nong  
108 Materia Medica (A. D. 102-200) of E. Han Dynasty and it was reported to be  
109 adopted to treat constipation and dysentery. Based on the modern  
110 investigations, the new application of Rhubarb involves cure of chronic renal  
111 failure, protection of damaged liver, and anti-aging effects (Xu et al. 2015; Lu  
112 et al. 2014; Xie and Sang 2014). *Rheum* (Polygonaceae) includes more than  
113 60 species distributed worldwide, with 39 species in the Western and Northern  
114 part of China, is one of the most important medicinal resources in China and  
115 one of the main plant source containing natural anthraquinone compounds

116 (Xiao et al. 1980). Among the anthraquinones reported to be present in  
117 Rhubarb (Ye et al. 2007), Sennosides (Sennoside A and Sennoside B) are  
118 considered to be the purgative components (Xiao et al. 1980; Zwaving 1972),  
119 Rhein and Physcion possess antitumor and anti-inflammatory properties  
120 (Zwaving 1972; He et al. 2011), with the latter presenting also  
121 hepatoprotective, antifungal and anticancer activities (He et al. 2011).  
122 Aloe-Emodin, Emodin and Chrysophanol have been reported to decrease the  
123 levels of serum total bilirubin, showing partial protective effects on cholestatic  
124 liver injury (Zhao et al. 2009). Glucosyl gallates (Maldonado et al. 2011),  
125 Naphthalenes (Tsuboi et al. 1977), and Catechins (Sill et al. 1974) exhibited  
126 potential antioxidant and anticancer properties. A number of polyphenol  
127 derivatives, including, for instance, anthraquinones, have been reported to  
128 interfere with the life cycle of different viruses, among which  
129 encephalomyocarditis virus in mice (Barnard et al. 1992), human  
130 cytomegalovirus (Barnard et al. 1992; Bamard et al. 1995), poliovirus (Semple  
131 et al. 2011) and hepatitis B virus (Dang et al. 2006), as well as to inhibit the  
132 catalytic activity of HCV NS5B Polymerase (Tramontano et al. 2011) and  
133 HIV-1 RT (Tramontano et al. 2011; Higuchi et al. 1991; Esposito et al. 2012;  
134 Esposito et al. 2011; Kharlamova et al. 2009) and integrase (IN) functions in  
135 biochemical assays (Tintori et al. 2015; Esposito et al. 2015). In particular, a  
136 number of anthraquinone derivatives were reported to affect both HIV-1 RDDP  
137 and RNase H RT-associated functions in biochemical assays, but were not

138 active on viral replication in cell-based assays (Tramontano et al. 2011;  
139 Higuchi et al. 1991; Esposito et al. 2012; Esposito et al. 2011; Kharlamova et  
140 al. 2009). Hence, we wanted to evaluate the effects of the chemical  
141 components originated from the Rheum Chinese plants to inhibit the HIV-1  
142 RT-associated and IN activities and to interfere with the HIV-1 life cycle in  
143 order to identify new scaffold inhibiting multiple HIV-1 targets.

144

## 145 **2. MATERIALS AND METHODS**

146 **2.1. Plant materials and reagents.** From June 2007 to September 2007, *R.*  
147 *palmatum* and *R. officinale* were collected from Gansu and Sichuan province  
148 respectively in China by one of the authors, and were naturally dried at room  
149 temperature. The raw materials were authenticated by Prof. Anren Li (Institute  
150 of Medicinal Plant Development, Beijing, China). The voucher specimens were  
151 deposited in the Laboratory of Pharmaphylogeny, Institute of Medicinal Plant  
152 Development (Beijing, China). The reference compounds Aloe-Emodin, Rhein,  
153 Emodin, Chrysophanol, Physcion were purchased from the National Institute  
154 for Control of Pharmaceuticals and Biological Products (Beijing, China), and  
155 Sennoside A and Sennoside B from the Mansite Pharmaceutical Co. Ltd.  
156 (Chengdu, China). HPLC grade acetonitrile and methanol were purchased  
157 from Burdick & Jackson (Muskegon, MI, USA). Deionized water was prepared  
158 by Milli-Q system (Millipore, Bedford, MA, USA). Phosphoric acid and

159 methanol were of analytical grade purchased from Beijing Beihai Fine  
160 Chemicals Co. Ltd. (Beijing, China).

161 **2.2 Preparation of sample solutions.** The dried roots of plant materials were  
162 grounded by an electric mill into powder (40 mesh). Individual samples (~200  
163 mg) were accurately weighed, then mixed with 30 ml methanol/water (30 ml,  
164 80:20, v/v) and sonicated for 60 min at room temperature. The extract solution  
165 was cooled down to room temperature, and appropriate amount of the solution  
166 was added to the original weight. The final solution was filtered through 0.22  
167 µm membrane and then injected 10 µl for each HPLC analysis (Fugure S1).

168 **2.3 Protein expression and purification.** HIV-1 RT gene subcloned into the  
169 p6HRT\_prot plasmid was kindly provided by Stuart Le Grice (NCI, Frederick,  
170 Maryland, USA). Protein expression and purification was performed in M15  
171 *Escherichia coli* cells as described (Esposito et al. 2012). Full-length IN and  
172 LEDGF proteins were expressed in BL21 *E. coli* (DE3) (Tintori et al. 2015;  
173 Esposito et al. 2015). HIV-1 K103N, Y108C and Y188L RT mutants were  
174 produced by site-directed mutagenesis using the Stratagene kit (Agilent  
175 Technologies, Milano, Italy), according to manufacturer's indication as  
176 described (Corona et al. 2014).

177 **2.4 RNase H polymerase-independent cleavage assay.** The HIV-1  
178 RT-associated RNase H activity was measured as previously described  
179 (Esposito et al. 2013).



180 **2.5 RDDP assay.** The HIV-1 RT-associated RDDP activity was measured  
181 using the Enz-Check Reverse Transcriptase Assay Kit (Life technologies,  
182 Carlsbad, California, USA), as previously described (Corona et al. 2014). The  
183 Yonetani-Theorell analysis was performed as previously reported (Esposito et  
184 al. 2011).

185 **2.6 Homogeneous Time Resolved Fluorescence (HTRF) LEDGF**  
186 **dependent and independent assay.** The IN LEDGF/p75 dependent assay  
187 allowed to measure the inhibition of the 3' processing and strand transfer IN  
188 reactions in the presence of recombinant LEDGF/p75 protein, as previously  
189 described (Tintori et al. 2015; Esposito et al. 2015).

190 **2.7 Cell line cultures.** The human T-lymphoblastoid Jurkat cell line, clone  
191 E6-1, and the human embryonic kidney 293T cell line were obtained from  
192 ATCC (Manassas, VA, USA) and maintained in RPMI 1640 or DMEM (GIBCO,  
193 Life Technologies, Monza, Italy), respectively, supplemented with 10%  
194 heat-inactivated fetal bovine serum at 37 °C in 5% CO<sub>2</sub> humidified  
195 atmosphere.

196 **2.8 Plasmids and Viruses.** HIV-1 stock was produced by transient  
197 transfection of Jurkat cells with the pSVC21 plasmid containing the infectious  
198 HXBc2 molecular clone of HIV-1 (Ratner et al. 1985) by the DEAE-dextran  
199 method as described previously (Cullen 1987) and stored at -80 °C until use.  
200 Viral titer was measured as 50% Tissue Culture Infective Doses (TCID<sub>50</sub>)/ml

201 on C8166 cells by the Reed and Muench end point dilution method (Reed and  
202 Muench 1983).

203 Recombinant HIV-1 HXBc2 CAT virus was produced by co-transfection of  
204 293T cells with the pSVC21 Vpr<sup>+</sup>Vpu<sup>+</sup>Nef<sup>+</sup>Δenv-CAT plasmid along with the  
205 pSVIIIenv plasmid (J. Sodroski, Harvard University, Boston), with the CaPO<sub>4</sub>  
206 method. The pSVC2.1 Vpr<sup>+</sup>Vpu<sup>+</sup>Nef<sup>+</sup>Δenv CAT plasmid (Sartori et al. 2011)  
207 was generated from the pSVC2.1 plasmid (J. Sodroski, Harvard University,  
208 Boston) which contains the complete proviral genome of the HIV-1 HXBc2  
209 (vif<sup>-</sup>-vpu<sup>-</sup>-vpr<sup>-</sup>-nef<sup>-</sup>) (Ratner et al. 1985), by inserting the Vpu, Vpr and Nef  
210 encoding sequences and deleting the BglIII-BglIII env region. The  
211 chloramphenicol acetyltransferase (CAT) reporter gene was inserted in the  
212 pSVC2.1 Vpr<sup>+</sup>Vpu<sup>+</sup>Nef<sup>+</sup>Δenv vector with resulting inactivation of the rev gene.  
213 The pSVIIIenv plasmid expresses the HIV-1 laboratory-adapted T-cell-tropic  
214 strain HXBc2 envelope glycoprotein along with Rev. Recombinant HIV-1  
215 virions expressing CAT were collected 48 hours post-transfection and filtered  
216 (0.45-µm-pore-size filter). Viral titer was determined in terms of Reverse  
217 Transcriptase (RT) activity (Rho et al. 1981).

218 **2.9 HIV-1 env complementation assay.** Jurkat cells (1 x 10<sup>6</sup>) were incubated  
219 with 30,000 <sup>3</sup>H cpm RT units of recombinant CAT reporter virus at 37°C in the  
220 absence or presence of different amount of compounds (5, 10 and 20 µM).  
221 Cells were lysed 4 days after infection and CAT activity was determined.  
222 Compound concentration required to inhibit early phases of HIV-1 replication

223 by 50% ( $EC_{50}$ ) was calculated by nonlinear regression analysis with Sigma  
224 Plot (Jandel Scientific). Only results within a linear range (HIV-1 LTR-driven  
225 reporter CAT gene expression, i.e. conversion of chloramphenicol to acetyl  
226 chloramphenicol above 50%) were elaborated.

227 **2.10 Cytotoxicity assay.** The cytotoxicity of the compounds on the Jurkat cell  
228 line was assessed using the MTT method (Mosmann 1993).

229 **2.11 Quantitative Real Time PCR Analysis of HIV-1 DNA.** Jurkat cells ( $1 \times$   
230  $10^6$ ) were infected with 30,000  $^3H$  cpm RT units of HIV-1 HXBc2 CAT virus in  
231 the absence or presence of drugs (Raltegravir was used at 0.5  $\mu M$ , Efavirenz at  
232 0.1  $\mu M$ , Sennoside A at 20  $\mu M$ , Sennoside B at 20  $\mu M$  and RDS1760 at 25  $\mu M$ ).  
233 Total intracellular DNA was purified from treated and untreated cells 16 hours  
234 post-infection with DNeasy blood and tissue kit (Qiagen, Limburg Netherlands),  
235 according to the manufacturer's instructions, and treated with 20 U/ml of DpnI  
236 for 2 hours at 37°C to remove contaminating plasmid DNA (Munir et al. 2013).  
237 The early (negative strand strong stop) and late (gag) reverse transcription  
238 products and the total HIV-1 DNA were measured using quantitative real-time  
239 PCR (qPCR), as previously described ( Yu et al. 2008; Doitsh et al. 2010;  
240 Cheney and McKnight 2010). Integrated HIV-1 DNA was detected by nested  
241 Alu-gag PCR, as reported (Yu et al. 2008; Liszewski et al. 2009), with some  
242 modifications. Pre-amplification reaction was performed using AmpliTaq Gold  
243 DNA Polymerase (Applied Biosystems, Monza, Italy) and consisted of one  
244 cycle of denaturation (10 min 95°C), 39 cycles of amplification (15 sec 94 °C,

245 30 sec 50 °C, 8 min 72 °C) and one cycle at 72 °C for 7 min. Quantitative PCR  
246 reactions were performed using TaqMan universal PCR master mix (Applied  
247 Biosystems ,Monza, Italy). The PCR consisted of one cycle of denaturation (10  
248 min at 95°C) followed by 50 cycles of amplification (15 s at 95°C, 1 min at 60°C)  
249 in an ABI Prism® 7000 Sequence Detection System (Applied Biosystems,  
250 Monza, Italy).

251 **2.12 Time of Addition.** Jurkat cells were infected with HIV-1 IIIB at a  
252 multiplicity of infection (M.O.I.) of 0.5 TCID<sub>50</sub>/cell. After 1 hour of incubation at  
253 37 °C, the cultures were washed twice and 10<sup>5</sup> cells were maintained in the  
254 absence or presence of drugs. Test and reference compounds were added at  
255 different time points after infection (0, 1, 2, 3, 4, 5, 6, 7, 8 hours). Virus  
256 production in the supernatant of infected cells was determined by p24 assay at  
257 31 hours post-infection, according to manufacturer's instructions (Innotest®HIV  
258 Antigen mAb, Fujirebio, Ghent, Belgium). Raltegravir was used at 1 µM,  
259 Efavirenz at 1.5 µM, Zidovudine at 0.4 µM, and Sennoside A at 250 µM.

260

### 261 **3. RESULTS AND DISCUSSION**

262 **3.1 Effect of extracts from Rheum Chinese species and of their chemical**  
263 **components of both HIV-1 RT-associated functions.** With the aim of  
264 identifying new scaffolds able to inhibit both HIV-1 RT-associated functions to  
265 be developed as novel antiviral drugs (Distinto et al. 2013; Distinto et al. 2013),  
266 we thought that TCM could be a valid source of compound biodiversity. Hence

267 we tested extracts from *R. palmatum* and *R. officinale*, firstly for their ability to  
268 inhibit the HIV-1 RNase H activity, for which no drugs are currently available  
269 (Corona et al. 2013). Results showed that extracts from both plants potently  
270 inhibited this enzyme function with IC<sub>50</sub> values of 0.9 and 0.25 µg/ml,  
271 respectively (Table S1).

272 Subsequently, single constituents of the extracts were purified by HPLC and  
273 seven phenolic components were found to be their major bioactive  
274 constituents (Table S1). Specifically, these components included  
275 anthraquinone derivatives such as Emodin, Crysophanol, Aloe-Emodin,  
276 Physcion, Rhein and Sennoside (both Sennoside A and Sennoside B). Hence,  
277 we investigated their ability to inhibit both HIV-1 RT-associated RNase H and  
278 RDDP functions using the RNase H selective diketo acid (DKA) derivative  
279 DS1759 (Corona et al. 2014) and the NNRTI Efavirenz as controls (Table 1).  
280 Physcion and Emodin were not significantly active on both HIV-1  
281 RT-associated RDDP and RNase H activities (as the maximum concentration  
282 tested they inhibited slightly less than 50% of enzyme activity), while  
283 Aloe-Emodine and Crysophanol inhibited both HIV-1 RT-associated activities  
284 with IC<sub>50</sub> values around 21-26 µM (Table 1). Differently, Rhein inhibited weakly  
285 the HIV-1 RT-associated RNase H function, but not its RDDP activity.  
286 Noteworthy, the most active components were Sennoside A and Sennoside B,  
287 that inhibited both HIV-1 RT-associated functions with IC<sub>50</sub> values in the 2-5  
288 µM range.

289 **3.2 Evaluation of the effects of Sennoside A and B on mutated RT.** Since  
290 Sennosides A and B were identified as novel dual functions RTIs, we wanted  
291 to compare them to known NNRTIs or RNase H inhibitors by using a series of  
292 previously described HIV-1 RT mutants. Thus, we firstly tested the effect of  
293 Sennoside A and B on three known single RT mutants involved in NNRTI  
294 resistance such as K103N, Y181C and Y188L RTs, using Efavirenz as positive  
295 control (Table 2). Results showed a significant difference between Sennoside  
296 A and B. In fact, in the presence of these mutations Sennoside A showed a 4-  
297 to 15-fold decrease in its potency of inhibition. On the contrary, Sennoside B  
298 exhibited a 2- to 4-fold decrease in the inhibitory activity on K103N and Y188L  
299 RTs, and displayed an  $IC_{50}$  value on Y181C RT similar to the one obtained on  
300 wild type RT (Table 2). Next, we tested the effects of Sennoside A and B on  
301 the RNase H activity of RTs bearing mutation in the 474 and 475 amino acid  
302 residues (N474A and Q475A RTs) that are highly conserved and were recently  
303 reported to be involved in the selective binding of the DKA derivative RDS1759  
304 (Kharlamova et al. 2009; Corona et al. 2014; Corona et al. 2014). Results  
305 indicated that, compared to wild type RT, the inhibitory activity of both  
306 compounds were affected by the presence of the two mutations, leading to a  
307 decrease in their efficacy by 5- to 12-fold (Table 2). Thus, in the case of RNase  
308 H mutants, Sennoside A and B displayed a similar reduction in the inhibitory  
309 activity.

310 Subsequently, considering the results obtained with Sennoside A on the  
311 NNRTI resistant RT mutants, we wanted to better characterize its behavior in  
312 the presence of Efavirenz in order to explore the possibility that the two  
313 compounds may interact with the same RT binding site. Therefore we tested  
314 the effect of increasing concentrations of Sennoside A and Efavirenz on the  
315 RT-associated RDDP activity and analyzed the data by a Yonetani-Theorell  
316 plot (Yonetani 1982) (Fig. 1). Such analysis can reveal whether simultaneous  
317 binding (or inhibition) of the HIV-1 RT enzyme by the two compounds is  
318 occurring or not. Results showed that Sennoside A and Efavirenz are  
319 kinetically mutually exclusive, suggesting that either they bind to the same site  
320 or the binding of one compound prevents the binding of the second one.

321 Taken together, this first set of data indicated that Sennoside A and B inhibit  
322 both RT-associated functions. In particular, Sennoside A showed a reduced  
323 ability to affect the RDDP activity of K103N, Y181C and Y188L RT mutants,  
324 similarly to Efavirenz, and kinetic analysis indicated that Sennoside A and  
325 Efavirenz are kinetically mutually exclusive. Overall, these results suggest that  
326 Sennoside A binding might involve the NNRTIs binding pocket, and thus inhibit  
327 the RT-associated RDDP function binding to this pocket. However, it is well  
328 known that Efavirenz and other NNRTIs increase RT-associated RNase H  
329 activity (Palanappian et al. 1995; Radzio and Sluis-Cremer 2008), while our  
330 biochemical studies showed that Sennoside A inhibits this function. The  
331 observation that Sennoside A is less effective in inhibiting the RNase H activity

332 in the presence of N474A and Q475A mutations, that were previously reported  
333 to strongly affect DKA derivatives binding and inhibitory activity (Corona et al.  
334 2014), suggests that Sennoside A could also recognize a second site that  
335 might be localized nearby or within RNase H domain. Overall, the binding of  
336 Sennoside A to the first site, in the polymerase domain, would lead to the  
337 inhibition of the HIV-1 RT-associated RDDP function, while the binding to the  
338 second site, in the RNase H domain, would lead to the inhibition of the HIV-1  
339 RNase H function.

### 340 **3.3 Evaluation of the effects of Sennoside A and B on HIV-1 IN activity.**

341 Since a number of DKA derivatives inhibiting the RNase H activity by binding  
342 to the RNase H domain were reported to inhibit the HIV-1 IN activity (Esposito  
343 et al. 2012; Esposito et al. 2011; Tintori et al. 2015; Esposito et al. 2015;  
344 Tramontano et al. 2005; Costi et al. 2013; Costi et al. 2014; Cuzzuculi Crucitti  
345 et al. 2014), we asked whether Sennoside A and B might affect also the IN  
346 catalytic function. Thus, we tested their ability to inhibit the HIV-1 IN strand  
347 transfer reaction in the presence of the LEDGF/p75 cellular cofactor, using  
348 Raltegravir as positive control (Table 2). Data showed that Sennoside A  
349 inhibited the HIV-1 IN strand transfer activity with an IC<sub>50</sub> value of 3.8 μM,  
350 while Sennoside B was 23-fold less potent.

### 351 **3.4 Pharmacophoric differences between Sennoside A and B.** Sennoside

352 A and B were shown to differently affect HIV-1 wild type and mutant RTs as  
353 well as IN strand transfer catalytic function. Hence, in order to investigate the



354 different behavior of the two isomers A (Threo) and B (Erythro), we calculated  
355 their global minimum energy conformation (Table S2). Sennoside A and B are  
356 optical stereoisomers and therefore they do not differ for their chemical  
357 properties; however their pharmacophoric features are diverse for their spatial  
358 orientation (Fig. 2). These overall sterical differences can explain the different  
359 performance of the two Sennoside stereoisomers. In this respect, considering  
360 the large molecular surface of both Sennoside molecules, the different  
361 localization of wide hydrophobic areas in the two stereoisomers can  
362 dramatically affect the potential establishment of hydrophobic bonds and  
363 therefore influence the biological activity.

364 **3.5 Characterization of the mechanism of action of Sennoside A and B in**  
365 **cell-based assays.** Given their ability to inhibit both the HIV-1 RT and IN  
366 functions in biochemical assays, we wanted to evaluate the effect of Sennoside  
367 A and B on the early phases of the HIV-1 replication. Thus, we selected a  
368 transient trans-complementation assay to assess the replicative potential of  
369 HIV-1 in a single round of infection using a lymphoblastoid T cell line as target  
370 (Helseth et al. 1990; Parolin et al. 2003). In this assay, an env-defective HIV-1  
371 HXBc2 Vpr<sup>+</sup>Vpu<sup>+</sup>Nef<sup>+</sup>Δenv provirus encoding the bacterial CAT gene was  
372 complemented by the envelope glycoprotein from the HXBc2  
373 laboratory-adapted T-tropic virus. The level of CAT expression in the cell  
374 lysates allowed us to examine early events in the infection process (Fig. 3).  
375 Results showed that Sennoside A inhibited HIV-1 replication with an EC<sub>50</sub>

376 value around 9  $\mu$ M in the absence of cytotoxicity, as determined by MTT assay,  
377 while Sennoside B was inactive at the highest tested concentration. In addition  
378 Crysophanol, even though it was able of inhibiting both HIV-1 RT-associated  
379 activities in biochemical assay (Table 1), did not exert any effect on HIV-1  
380 replication.

381 In order to identify the viral process targeted by Sennoside A, we examined its  
382 effect on both reverse transcription and integration process by quantitative real  
383 time PCR (qPCR) analyzing viral cDNA intermediates in Jurkat infected cells  
384 treated with the drug. The primers and probe set used were selected in order  
385 to monitor sequential steps within the reverse transcription process,  
386 corresponding to generation of negative strong-stop DNA (early) and DNA  
387 strand elongation (late), and to quantify viral DNA that have successfully  
388 completed the integration process (Fig. 4A). Moreover, we assessed total viral  
389 DNA synthesis, including reverse transcription products that have completed  
390 the second strand transfer, as well as linear and circular form of viral DNA and  
391 integrated DNA (Fig 4A). Compounds were added at the time of infection and  
392 DNA collected at 16 hours post-infection. Efavirenz, Raltegravir and the DKA  
393 derivative RDS1760 (Corona et al. 2014) were used as positive controls (Fig.  
394 4B). Results showed that negative strong stop DNA synthesis was only slightly  
395 inhibited by Sennoside A at 20  $\mu$ M concentration (Fig. 4B). In contrast, late  
396 reverse transcription products generated after the first strand transfer  
397 decrease in the presence of Sennoside A, but not in the presence of

398 Sennoside B. The reduction of total viral DNA synthesis further confirmed this  
399 result.

400 Based on these observations, Sennoside A seemed to exhibit a mechanism of  
401 action similar to the one displayed by Efavirenz, affecting the RT activity during  
402 or soon after the first template exchange. Integration assay with an Alu-gag  
403 qPCR revealed that treatment with Sennoside A reduced also viral integration.  
404 Next, in order to gain further insights into the mechanism of action of  
405 Sennoside A, we performed a time of addition study in which compounds were  
406 added at the time of infection and every hour up to 8 hours after infection,  
407 using Efavirenz, Zidovudine and Raltegravir as controls (Fig. 5). Jurkat cells  
408 were infected with HIV-1 (Pauwels et al. 2011) and compounds were added  
409 with time lags of up to 8 hours after infection. According to the target of drug  
410 action or to the stage of interaction, addition of compound could be delayed for  
411 a specific number of hours without loss of antiviral activity. Results showed  
412 that Sennoside A profile of inhibition was similar to the one observed for the  
413 nucleoside analog Zidovudine (Fig. 5). These data confirm RT as a target of  
414 interaction of Sennoside A. On the other hand, our experimental setting cannot  
415 exclude that Sennoside A might have a slight effect also on HIV-1 IN.

416

#### 417 **4. CONCLUSIONS**

418 Starting from the TCM heredity and knowledge, we successfully identified in  
419 the Rhubarb the chemical component Sennoside A that is able to inhibit in the

420 low micromolar range both HIV-1 RT-associated activities, IN function and  
421 virus replication. Mode of action studies performed in cell cultures,  
422 demonstrated that Sennoside A targets the HIV-1 reverse transcription  
423 process, while it seems to be unable to significantly affect IN activity. Overall,  
424 Sennoside A represents a novel attractive scaffold of dual function RTI that  
425 deserves further investigations by means of chemical modification, in search of  
426 new dual enzyme derivatives active on both HIV-1 RT and IN.

427

#### 428 **Acknowledgments**

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430 Superiore di Sanità grant n. 40H98, and by 60% grant from the University of  
431 Padova.

432

#### 433 **Conflict of Interest**

434 We wish to confirm that there are no known conflicts of interest associated with  
435 this publication and there has been no significant financial support for this work  
436 that could have influenced its outcome.

437

438

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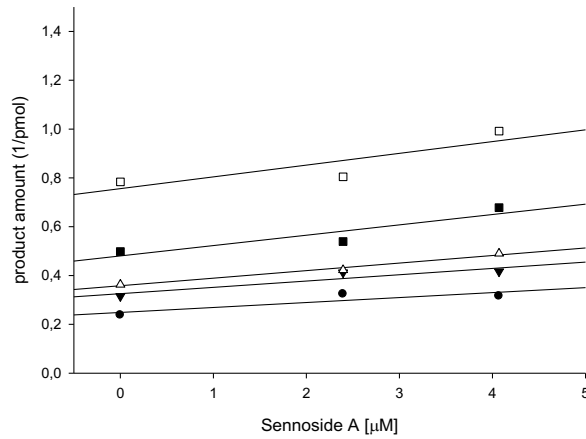
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611 **Figure 1. Yonetani–Theorell plot of the combination of Sennoside A and**  
612 **Efavirenz on the HIV-1 RT RNA-dependent DNA polymerase activity.**

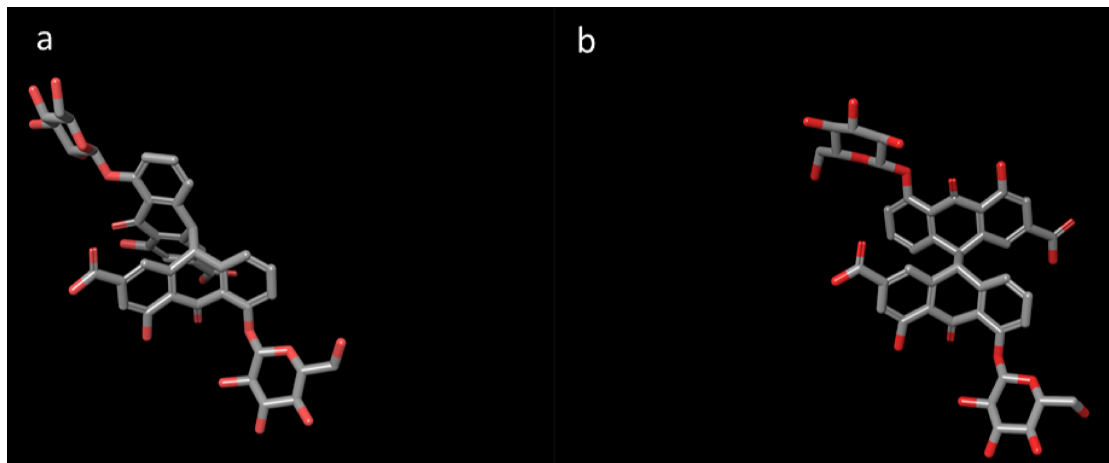
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615 HIV-1 RT was incubated in the presence of Sennoside A alone (●) or in  
616 the presence of different Efavirenz concentrations: 6 nM (▼), 9 nM (Δ), 13  
617 nM (■), 20 nM (□). In Y axis it was represented the reverse of the  
618 product formation.

619 **Figure 2.**



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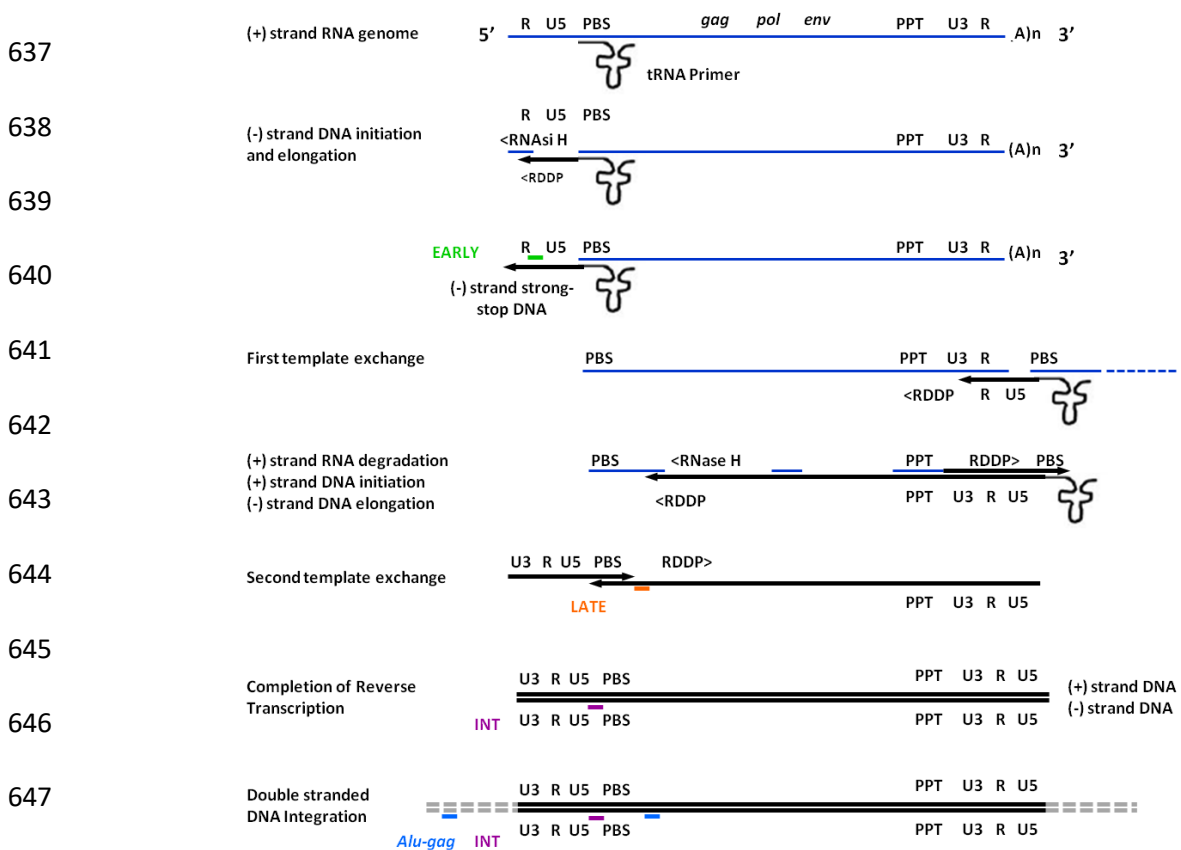
621 **Figure 3. Effect of Sennoside A and B on HIV-1 replication.**

622 **Figure 4. Quantitative determination of HIV-1 reverse transcription and integration**  
623 **products in the presence of Sennoside A and B.** Quantitative Real Time PCR analysis was  
624 used to measure the amount of HIV-1 early and late reverse transcription products,  
625 integrated and total DNA in the presence of test and reference compounds. **A.** Schematic

626 representation of primer set positions within reverse transcription and integration  
 627 intermediates. **B.** Total intracellular DNA was extracted from Jurkat infected cells 16 hours  
 628 post infection. Test and reference compounds were added at the time of infection (Efavirenz  
 629 0.1  $\mu$ M, Raltegravir 0.5  $\mu$ M, Sennoside A 20  $\mu$ M, Sennoside B 20  $\mu$ M, RDS1760 25  $\mu$ M). The  
 630 level of each viral DNA products was calculated as fold change relative to infected untreated  
 631 cells using  $\Delta\Delta$ Ct method (Livak and Schmittgen 2001). The same positive control (no drug)  
 632 was used for all analyses, thereby allowing the effect of Sennoside A and B on reverse  
 633 transcription and integration processes to be evaluated.  $\beta$ -globin housekeeping gene was  
 634 used as endogenous control. Values are means of duplicate determinations.

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636 **A**



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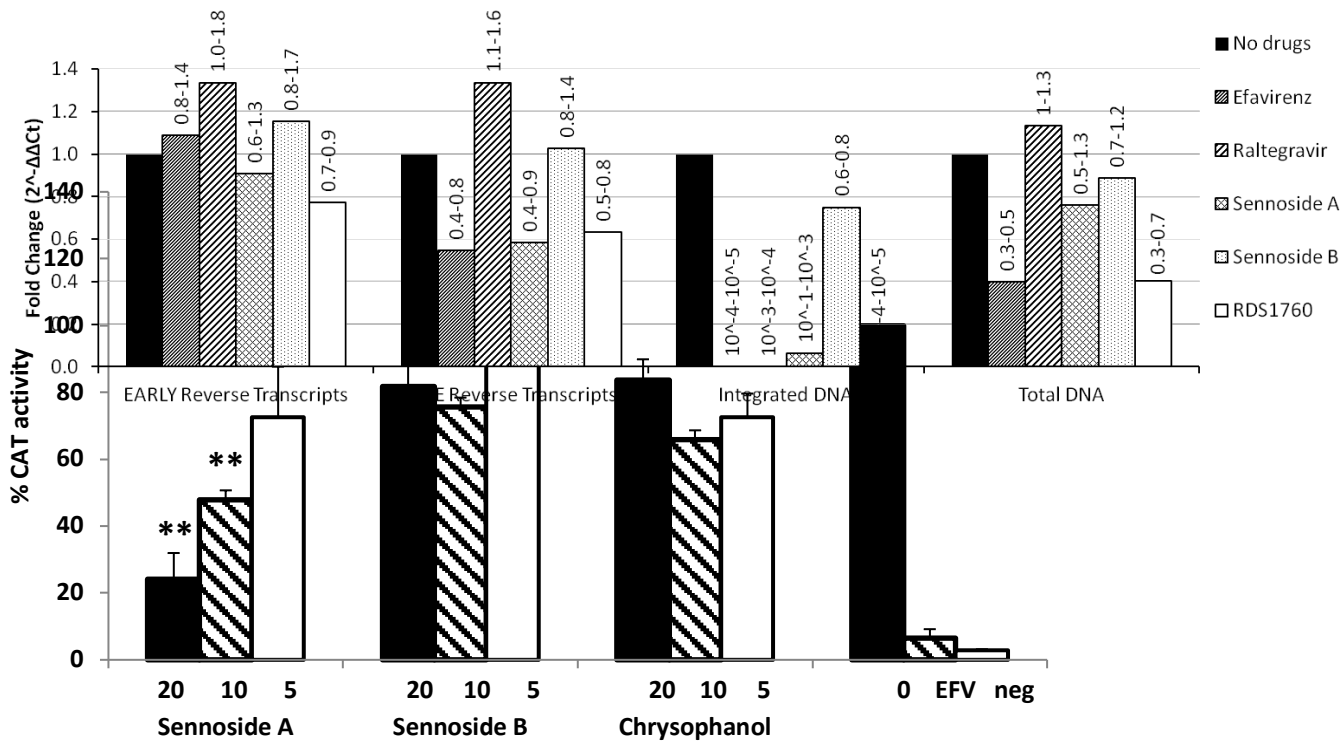
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658 **B**

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663 CAT activity was determined in Jurkat cells infected with HIV-1 recombinant CAT virus  
 664 pseudotyped with the envelope glycoprotein from the HXBc2 laboratory-adapted T-tropic  
 665 virus, in the absence (0) or presence of 20 μM, 10 μM and 5 μM of selected compounds 72  
 666 hours post infection. The results are reported as percent of conversion of  
 667 [<sup>14</sup>C]chloramphenicol to acetylated forms relative to the positive control (0). neg, negative

668 control; EFV, efavirenz. The means of two independent experiments and standard deviations  
669 are presented. \*\* indicates p=0.01.

670 **Figure 5. Time of addition.** Jurkat cells were infected with HIV-1 and the test  
671 compounds were added at different time points after infection. Virus production  
672 was determined by p24 Ag production in the supernatant at 31 hours post  
673 infection (h.p.i.). No drugs (positive control, ■); Raltegravir, 1  $\mu$ M, (x); Efavirenz,  
674 1.5  $\mu$ M ( $\blacktriangle$ ); Zidovudine, 0.4  $\mu$ M ( $\bullet$ ); Sennoside A, 250  $\mu$ M (o). This graph is  
675 representative of data from 2 independent experiments.

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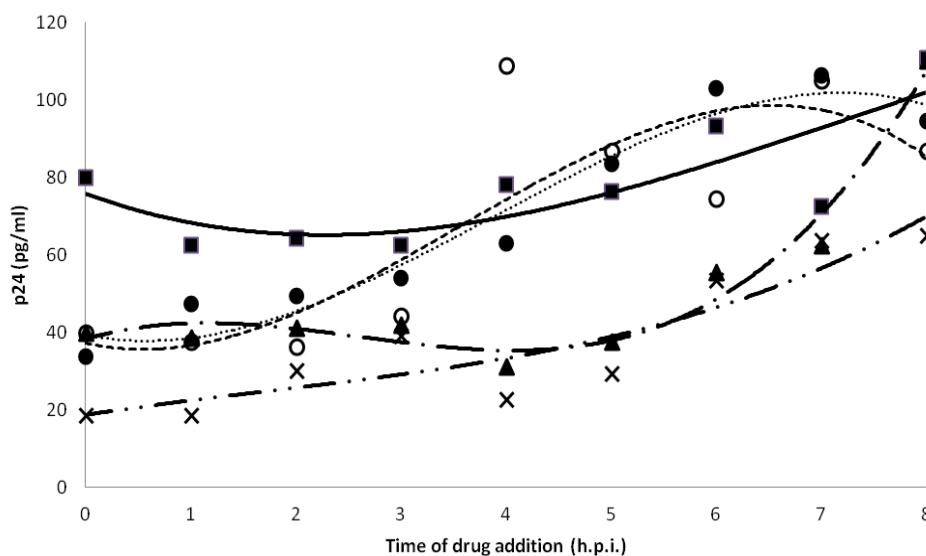
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695 **Table 1. Description of Species, collecting place, origin and components of Rheum**  
 696 **Chinese and their effects on HIV-1 RNase H.**

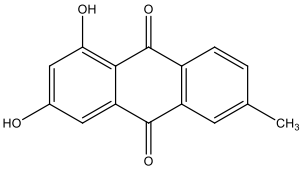
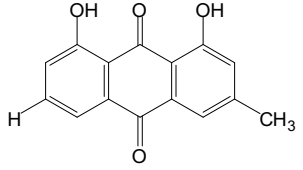
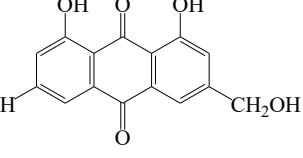
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Species	Place of collection	Origin	Aloe Emodin (mg/g)	Rhein (mg/g)	Emodin (mg/g)	Chrysophanol (mg/g)	Physcion (mg/g)	Sennoside A (mg/g)	Sennoside B (mg/g)	H RN a (μ
<i>R. palmatum</i> (D-13)	Tianzhu, Gansu	Wild	0.023	-	0.046	0.055	0.102	0.864	0.489	
<i>R. officinale</i> (D-15)	Songpan, Sichuan	Wild	0.05	0.039	0.033	0.107	0.16	0.162	0.037	0

698 Extract concentration required to inhibit the HIV-1 RT-associated RNase H activity by 50%

699

700 **Table 2. Effect of Rheum single components on the HIV-1 RT-associated**  
 701 **activities.**

Compound	Chemical structure	<sup>a</sup> IC <sub>50</sub> (μM)	
		RNase H	RDDP
<b>Emodin</b>		>100 (59%) <sup>b</sup>	>100 (51%)
<b>Chrysophanol</b>		25.0 ± 4.0	26.5 ± 2.5
<b>Aloe-Emodin</b>		23.0 ± 5.0	21.0 ± 4.0

<b>Physcion</b>		> 100 (53%)	>100 (57%)
<b>Rhein</b>		60 ± 12	>100 (62%)
<b>Sennoside A</b>		1.9 ± 0.3	5.3 ± 0.1
<b>Sennoside B</b>		2.1 ± 0.2	2.3 ± 0.4
<b>Efavirenz</b>		> 50 <sup>c</sup>	0.025 ± 0.0005 <sup>c</sup>
<b>RDS1759</b>		7.4 ± 0.2 <sup>c</sup>	> 50 <sup>c</sup>

702 <sup>a</sup>Compound concentration required to reduce HIV-1 RT-associated RNase  
703 H and RT-associated RNA Dependant DNA Polymerase activities by  
704 50%.

705 <sup>b</sup>Percentage of control activity in the presence of 100 μM concentration of  
706 compound.

707 <sup>c</sup>Data were expressed in μM.

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709